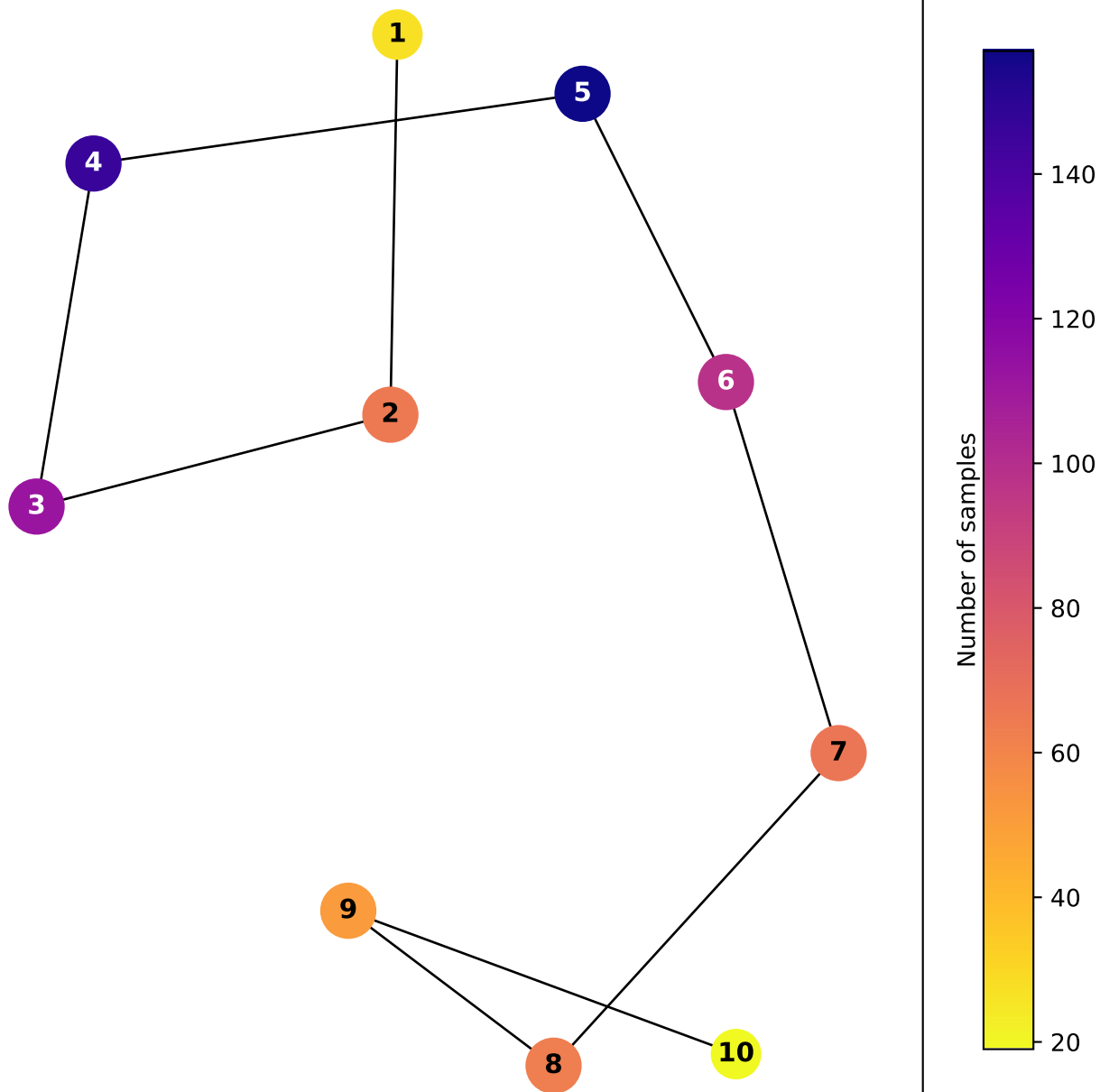


Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

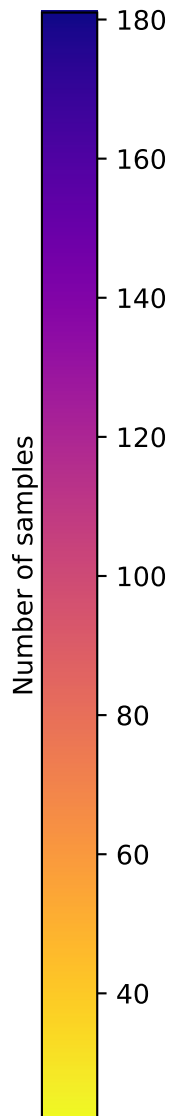
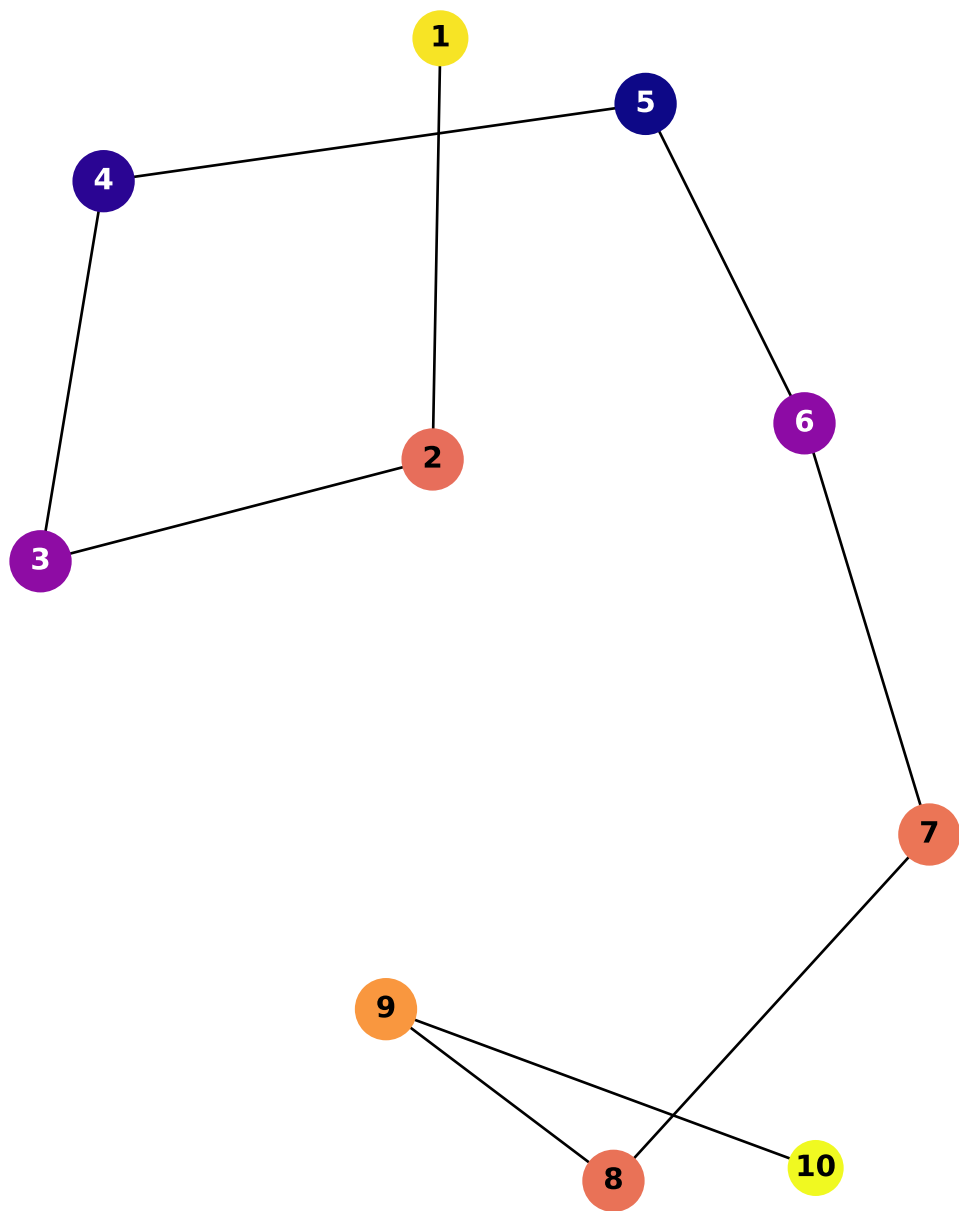
Intervals: 10 - Overlap: 0.1



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

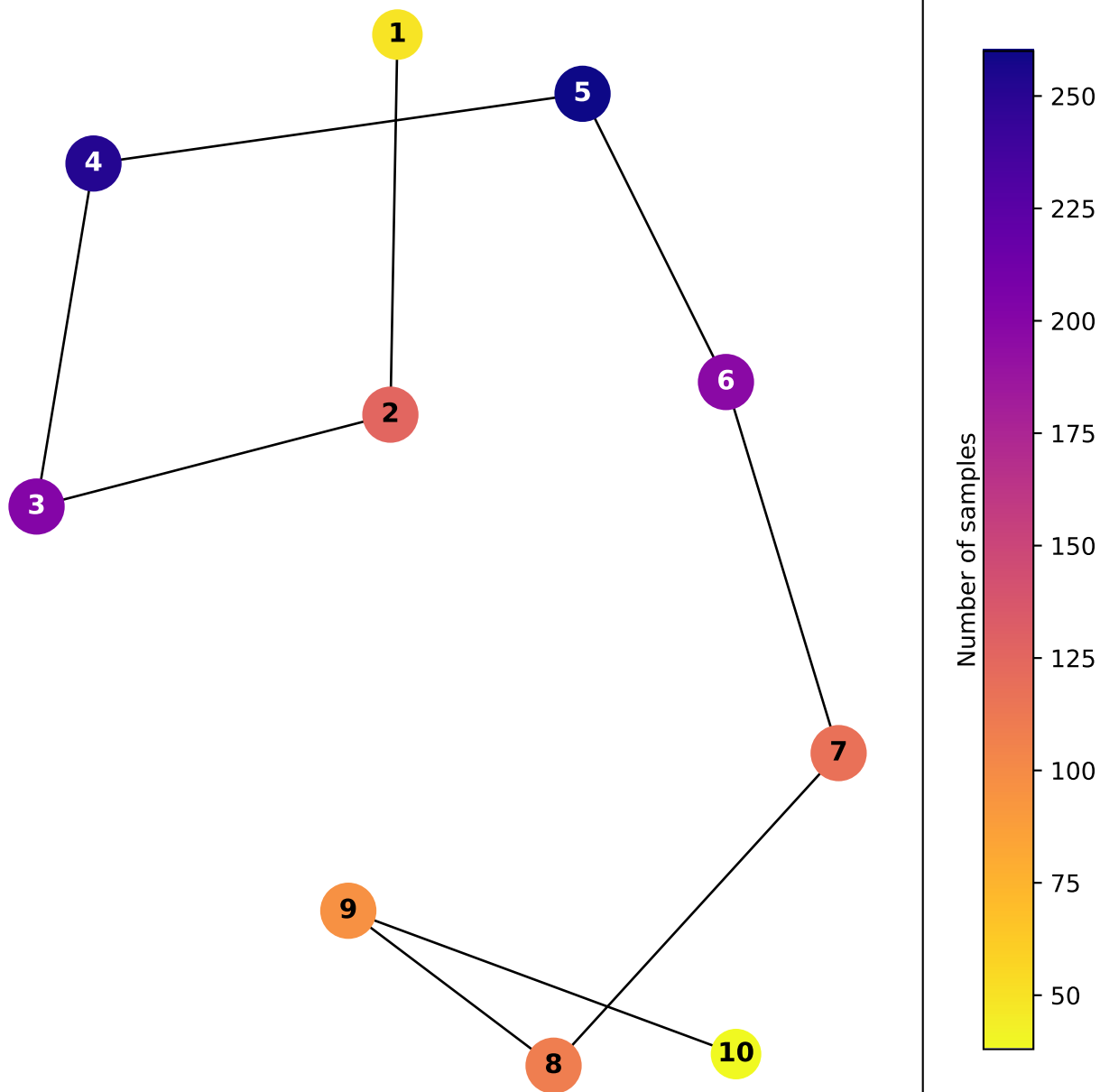
Intervals: 10 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

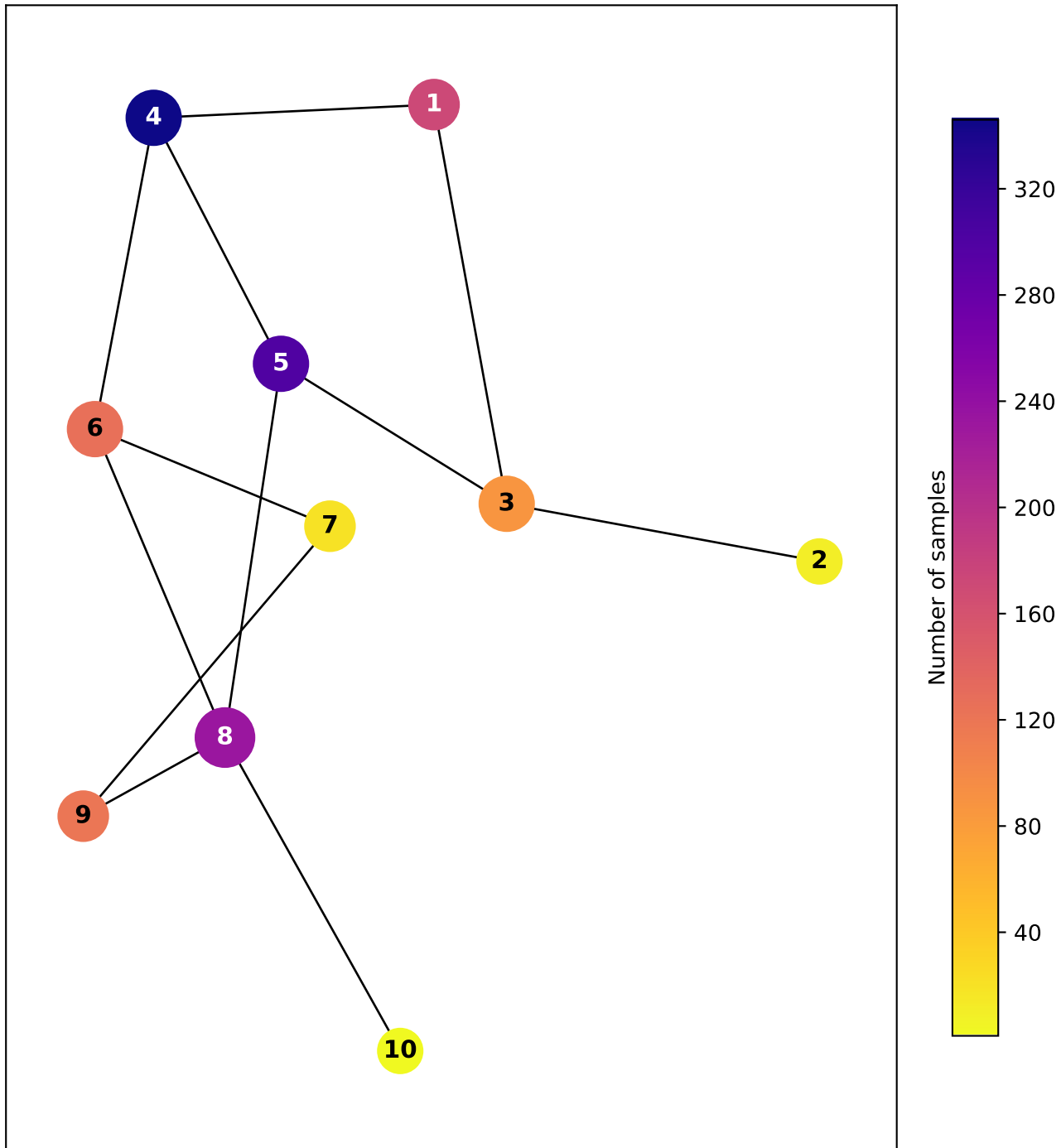
Intervals: 10 - Overlap: 0.5



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=2)

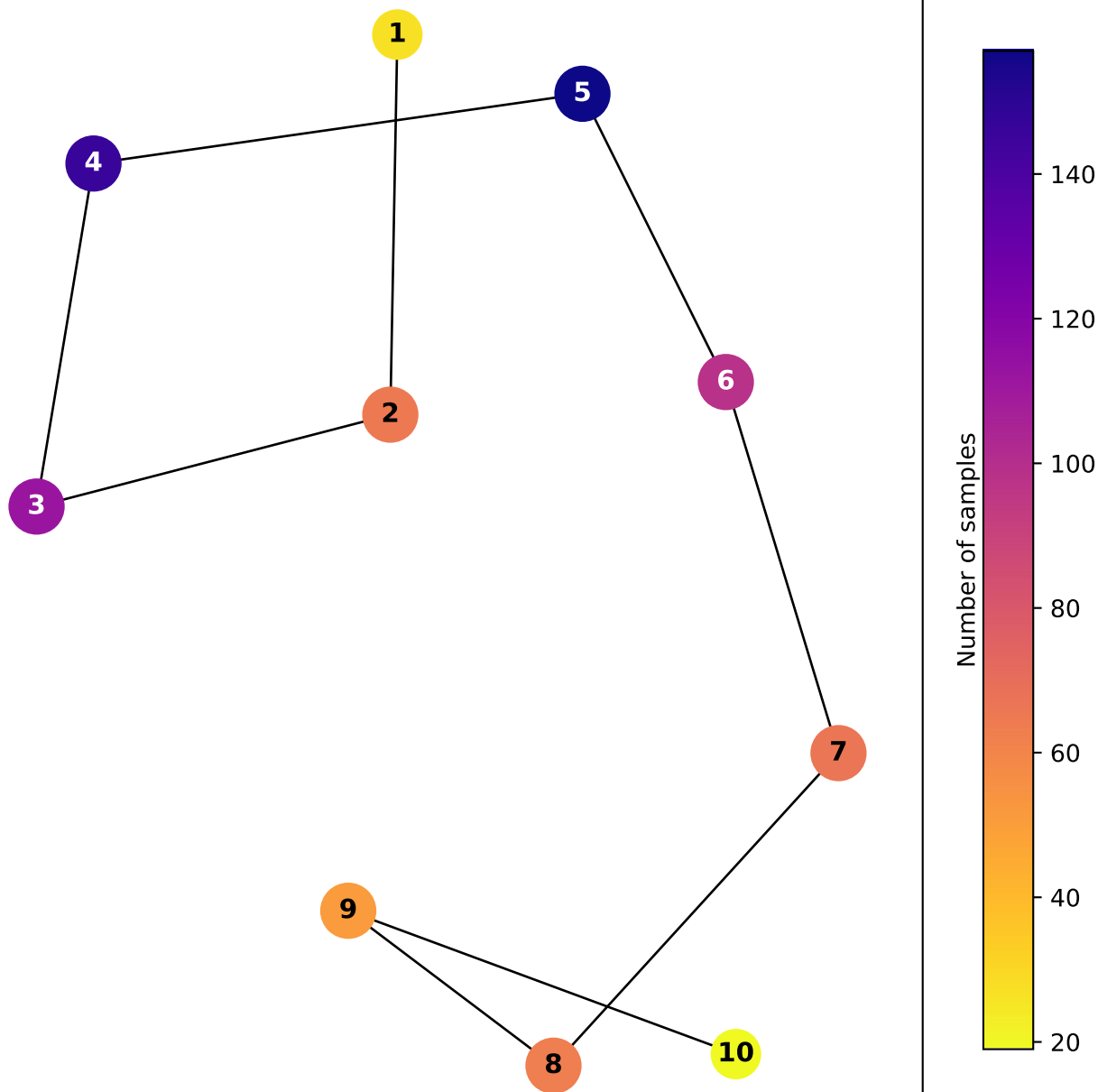
Intervals: 5 - Overlap: 0.5



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

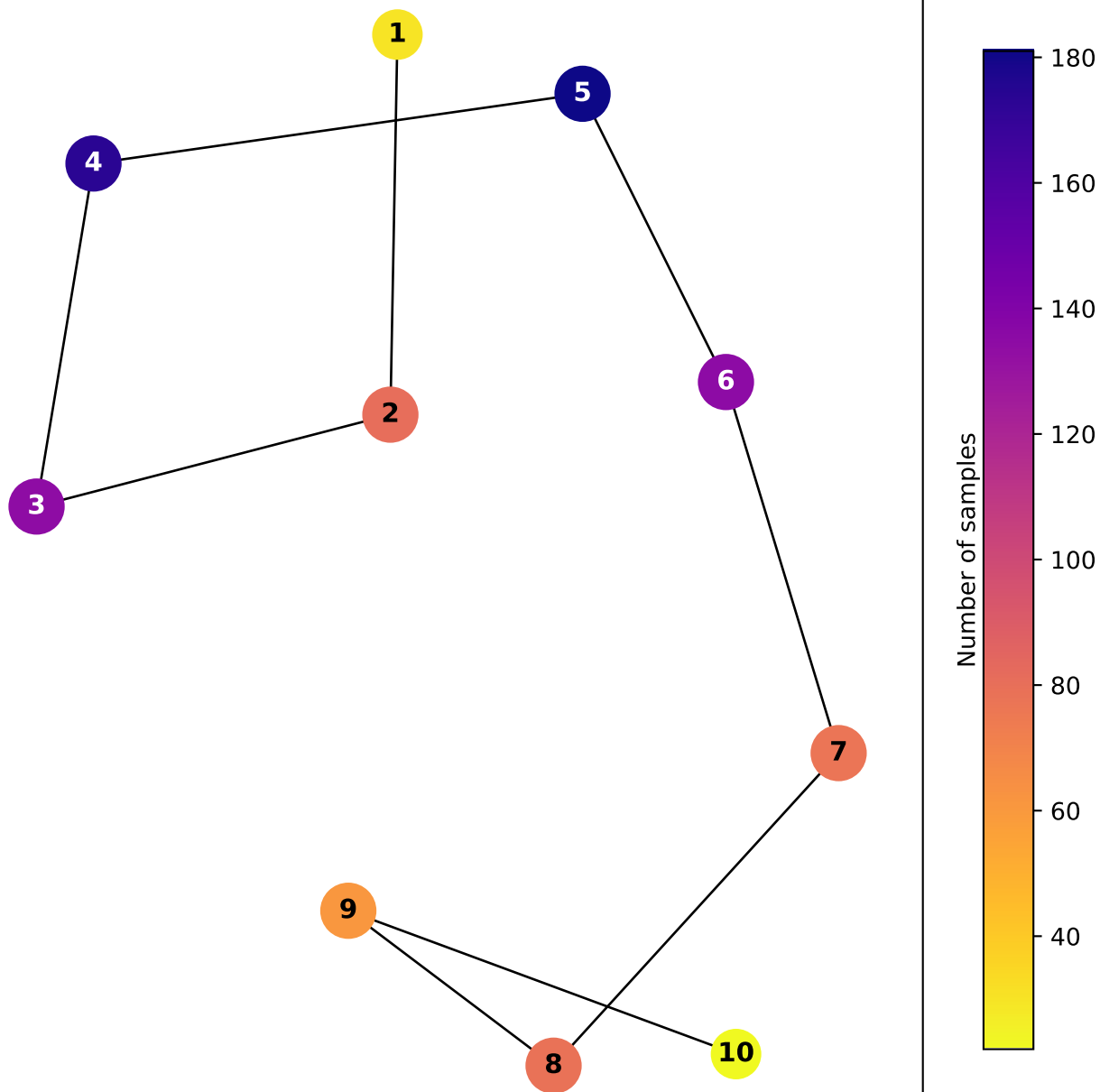
Intervals: 10 - Overlap: 0.1



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

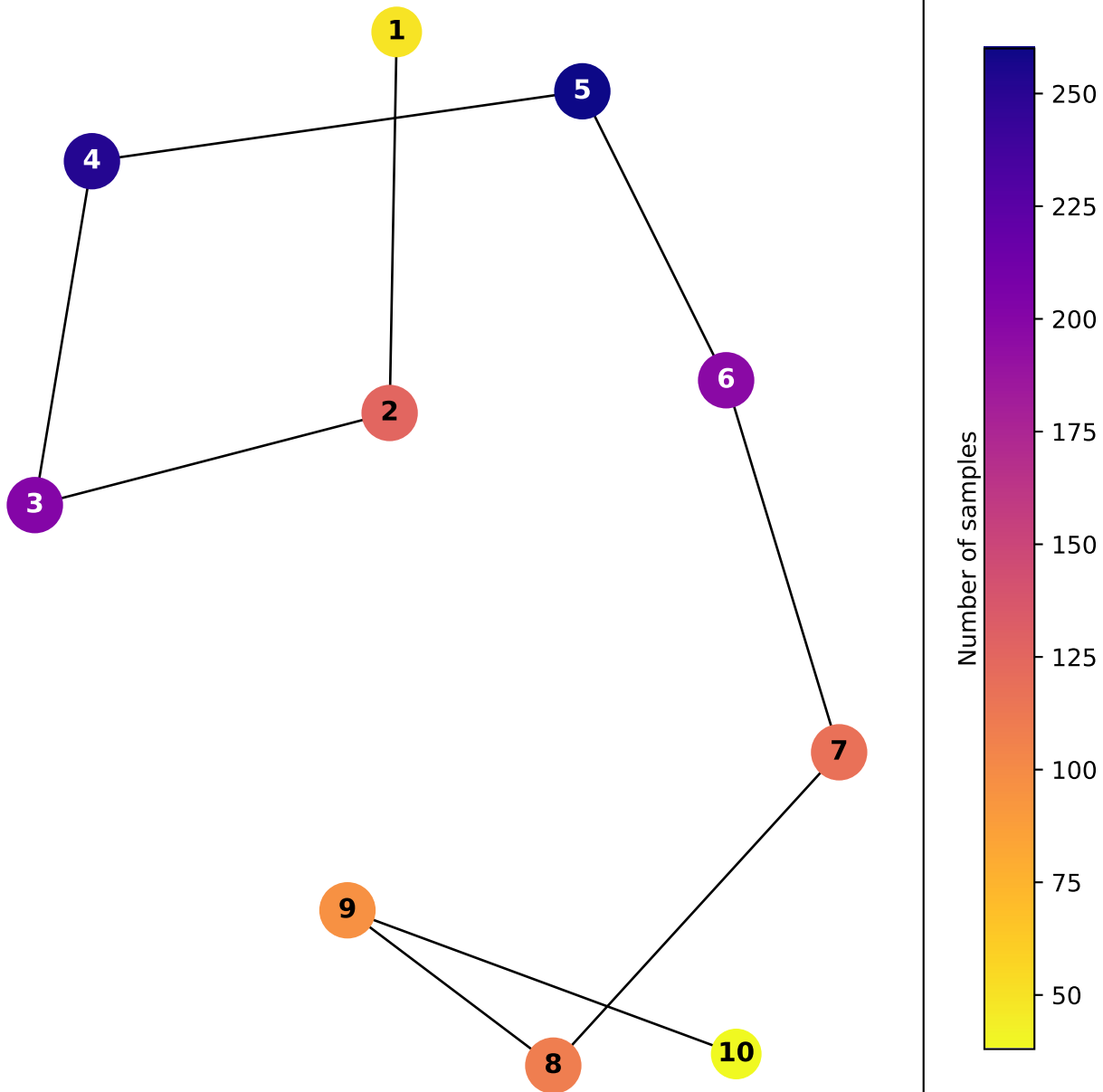
Intervals: 10 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

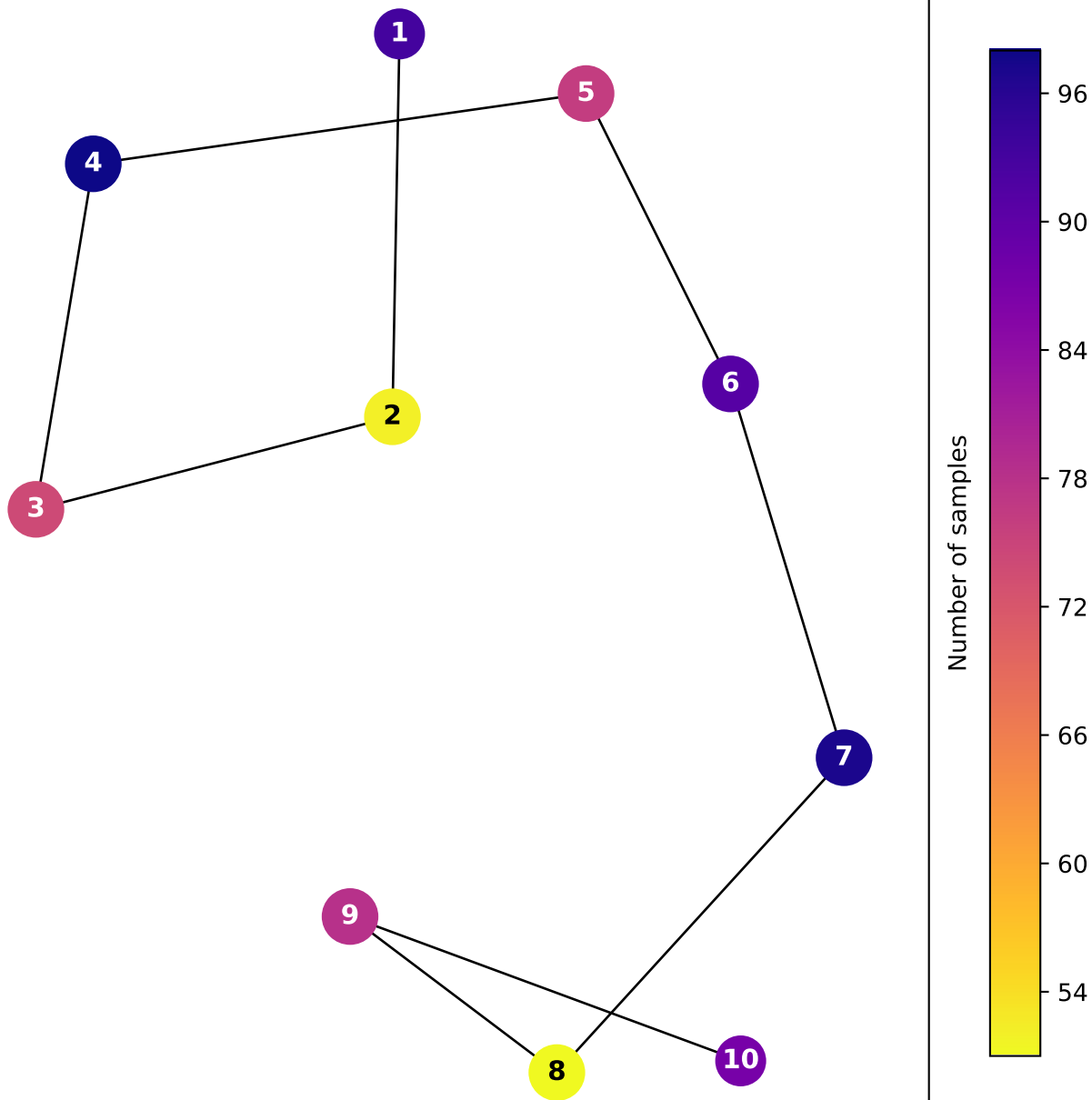
Intervals: 10 - Overlap: 0.5



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

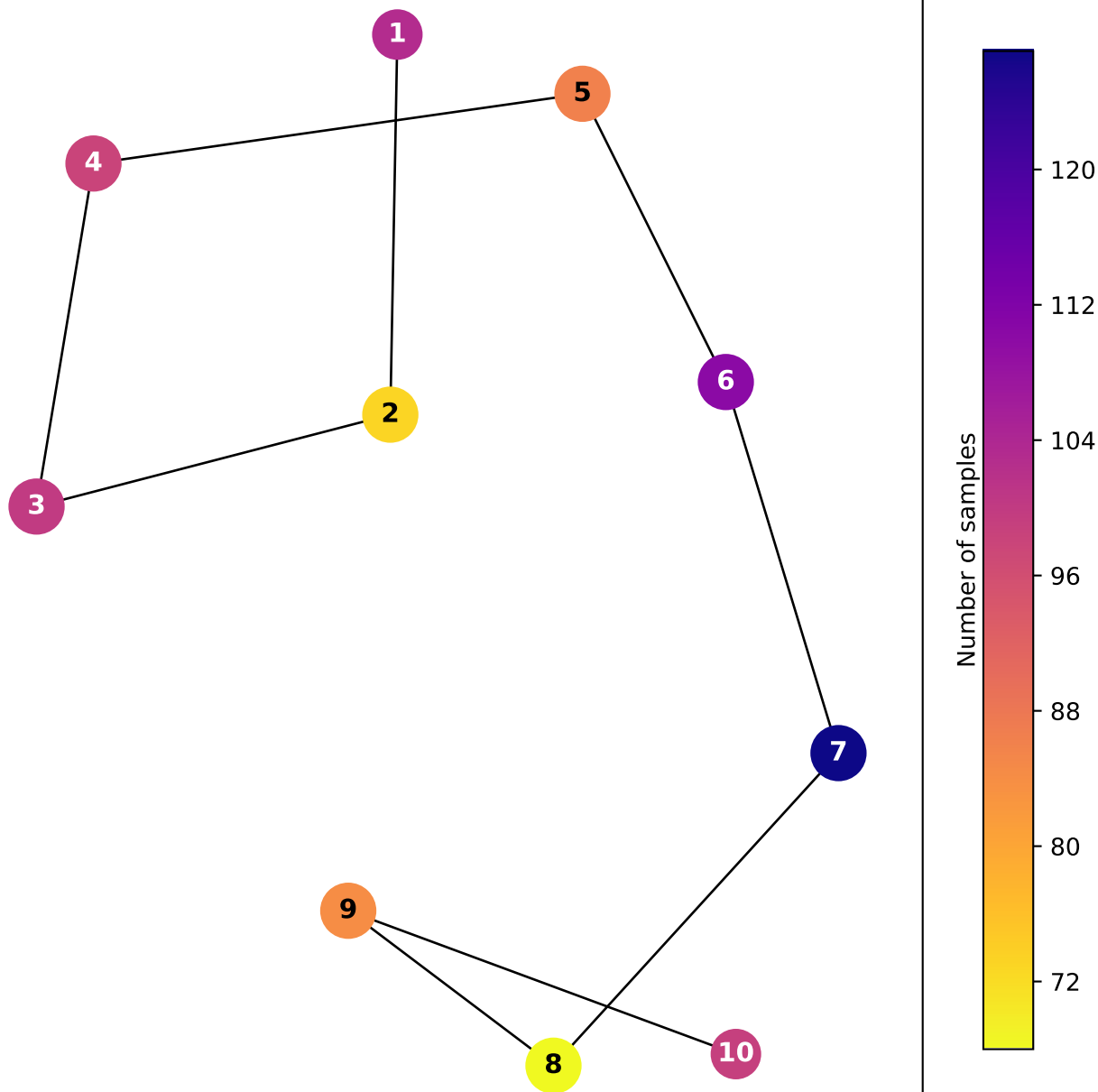
Intervals: 10 - Overlap: 0.1



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

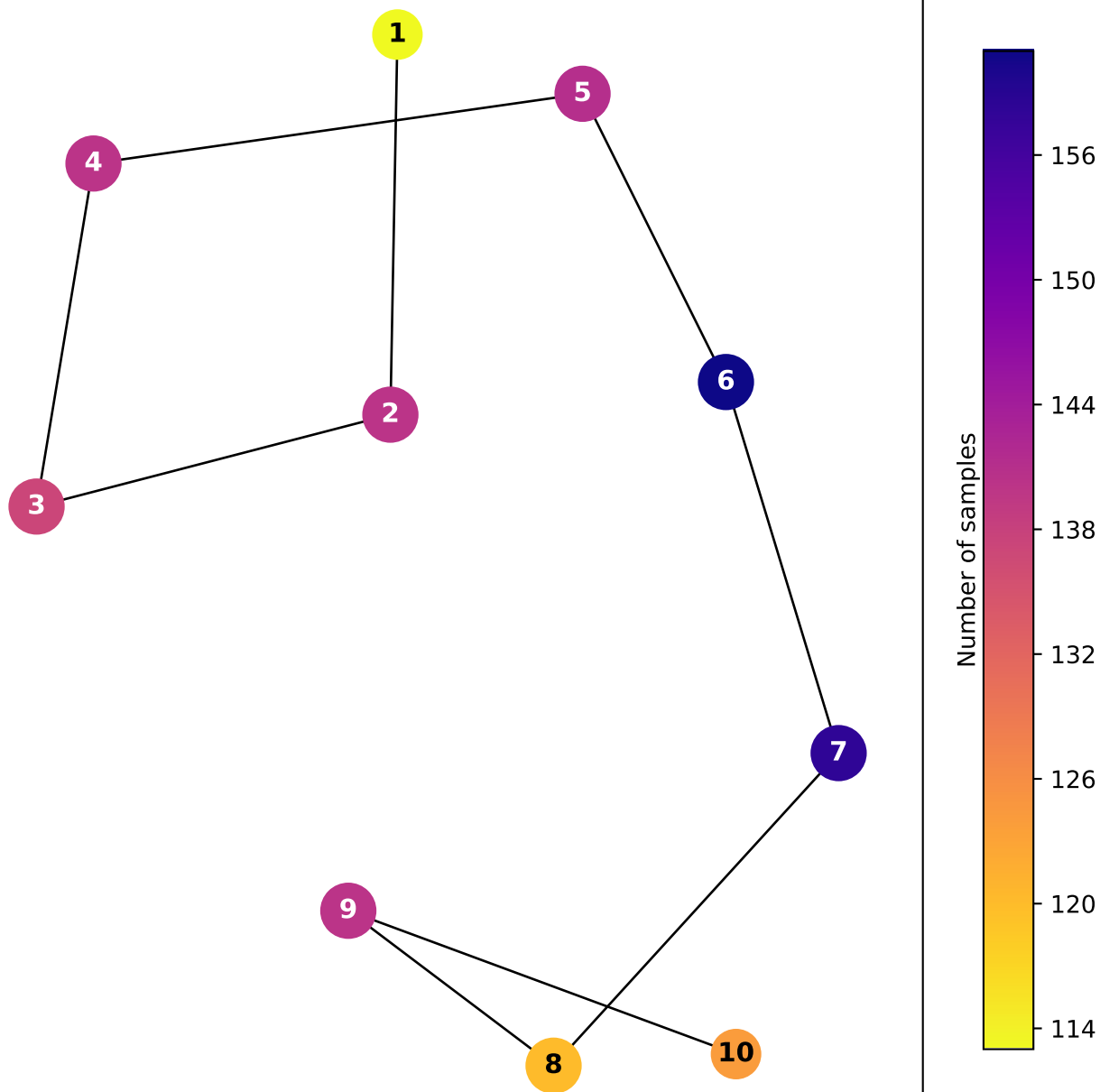
Intervals: 10 - Overlap: 0.25



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

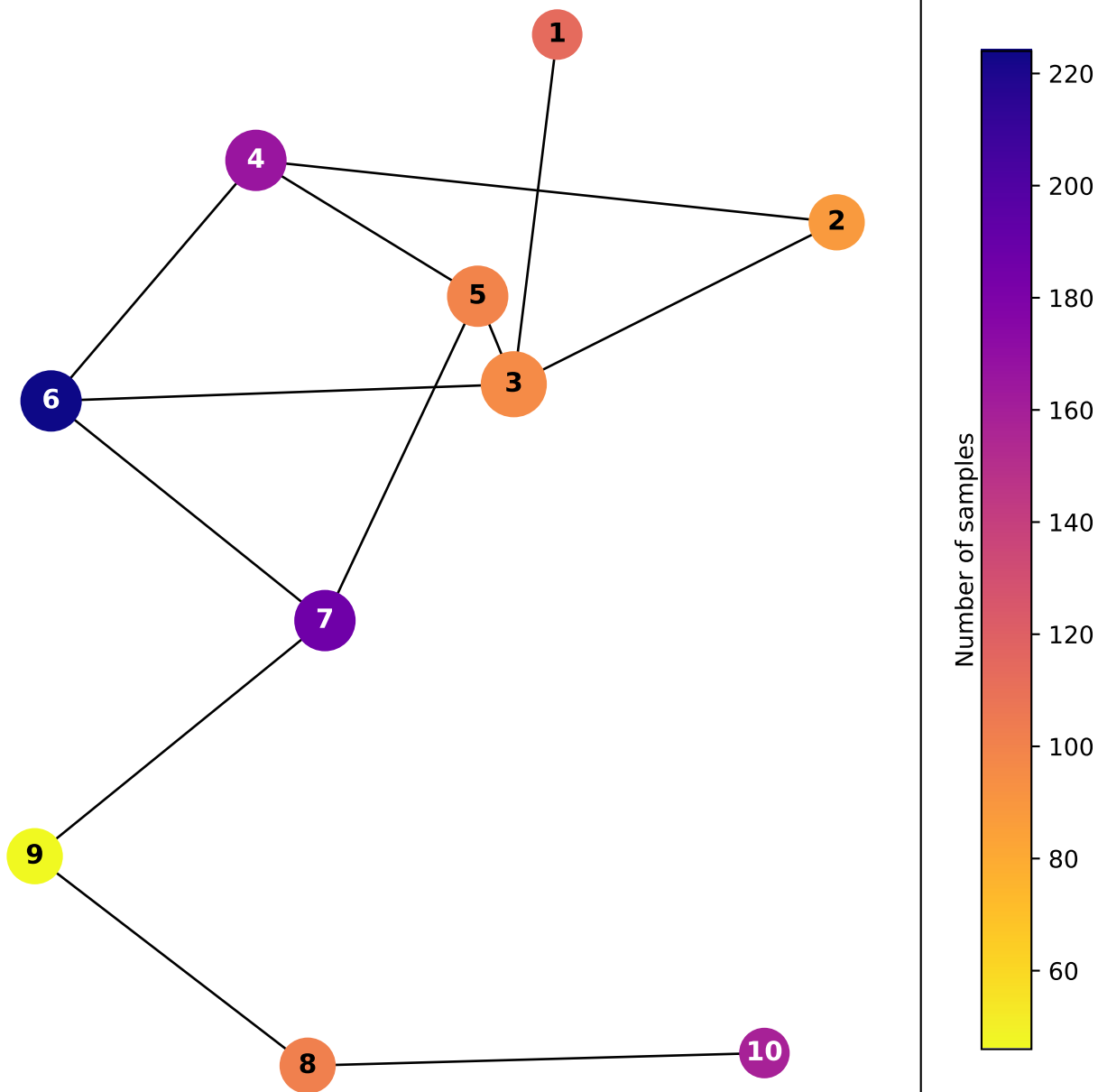
Intervals: 10 - Overlap: 0.5



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=2)

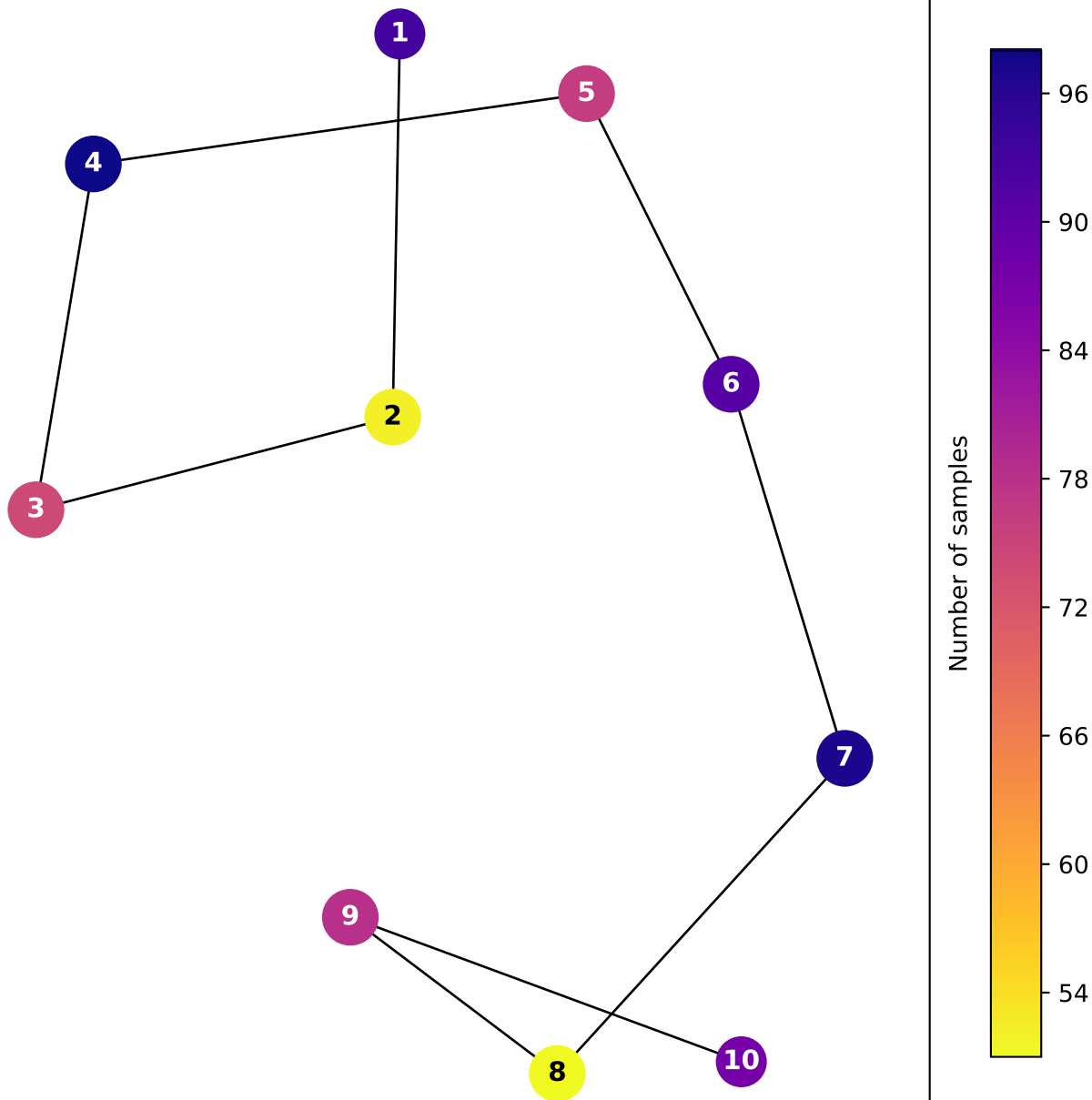
Intervals: 5 - Overlap: 0.5



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

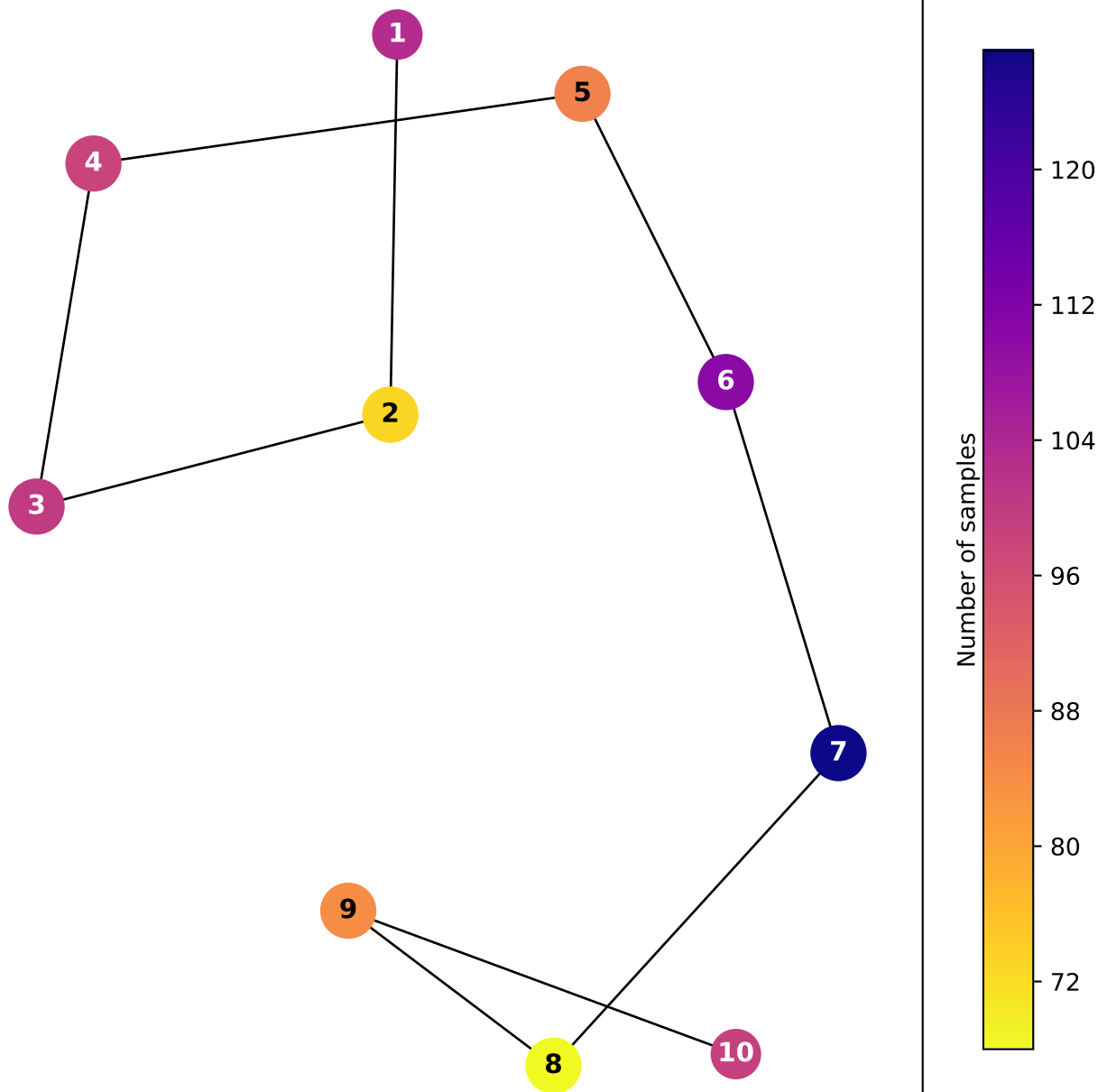
Intervals: 10 - Overlap: 0.1



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

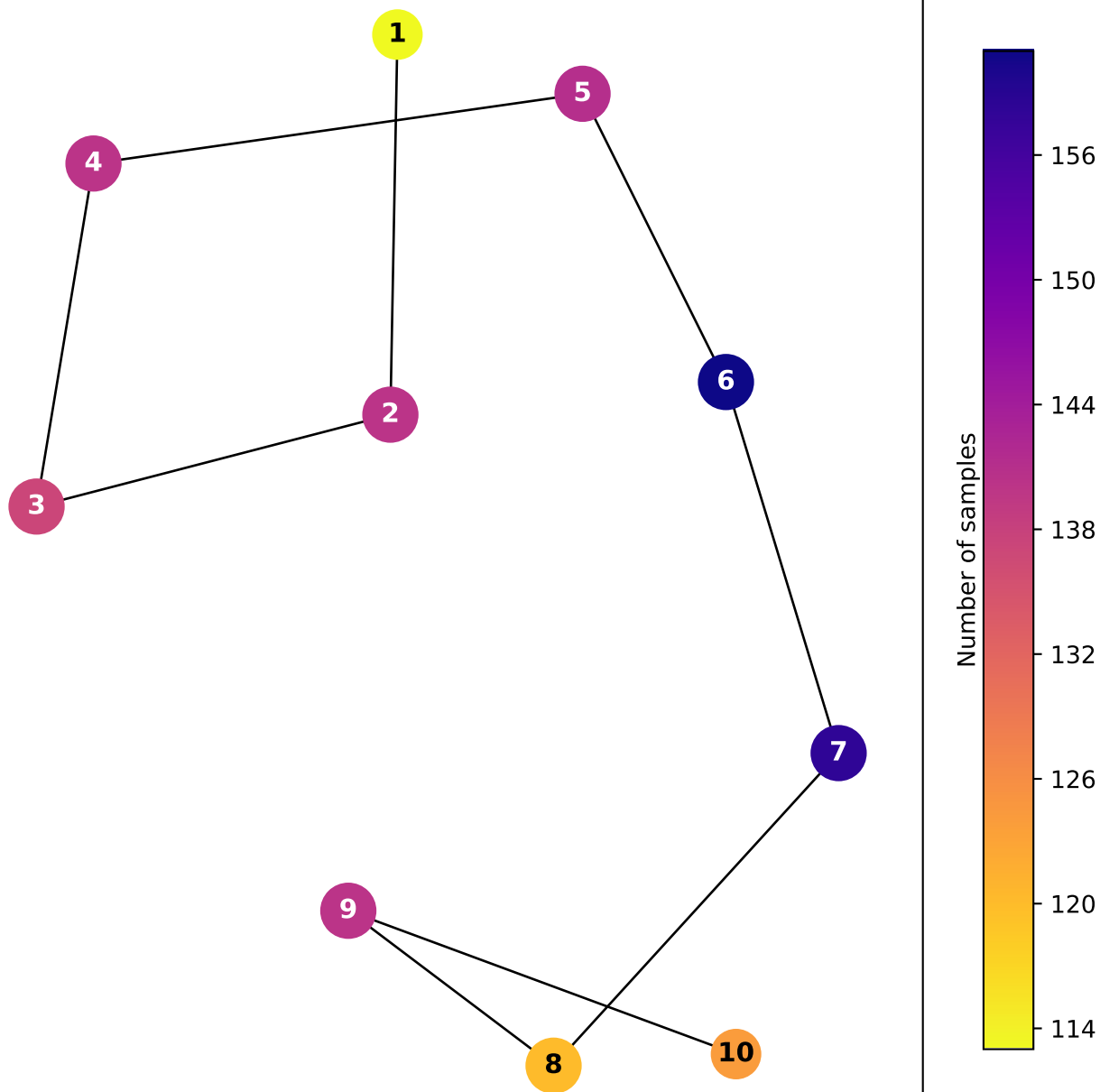
Intervals: 10 - Overlap: 0.25



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

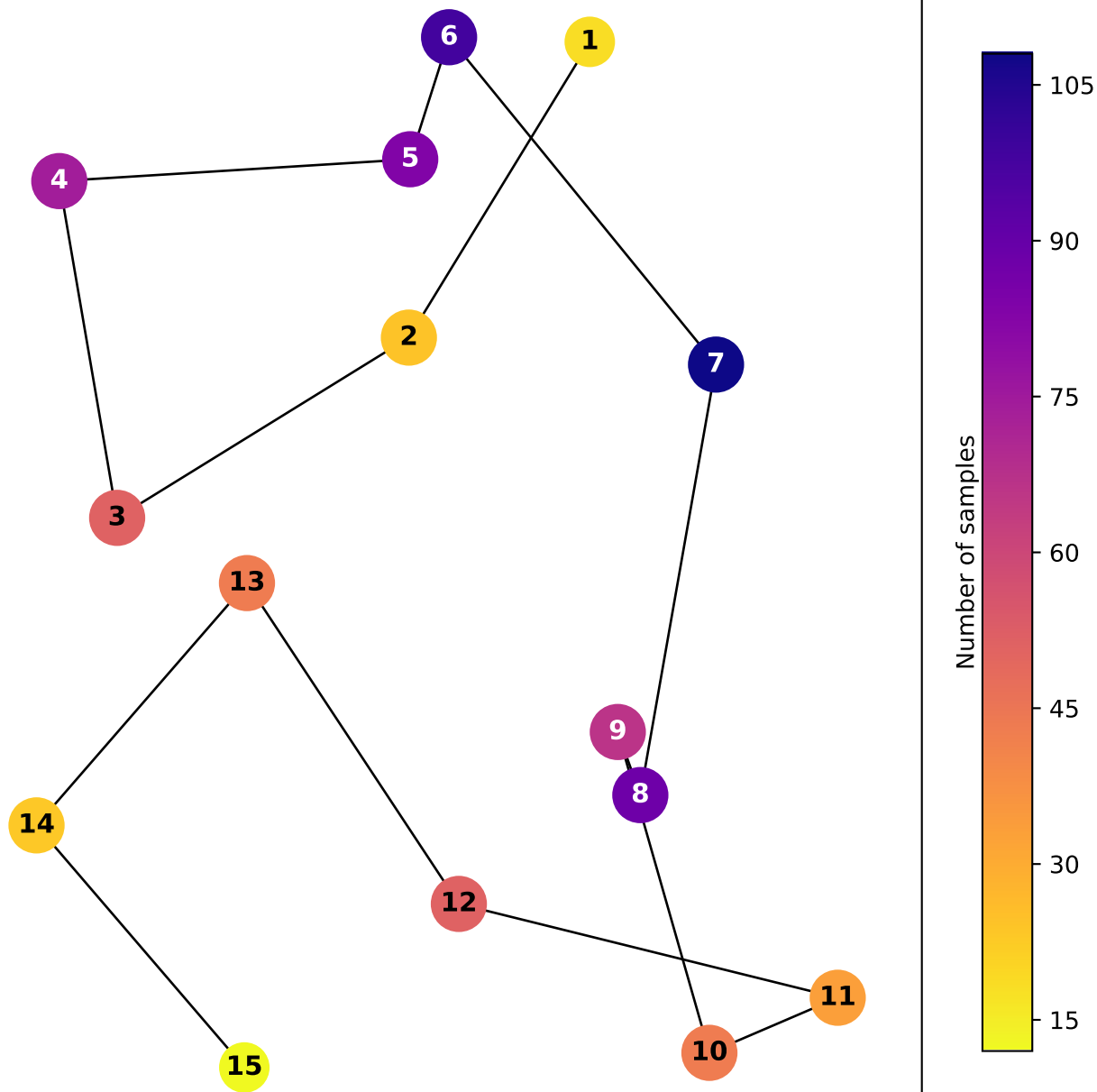
Intervals: 10 - Overlap: 0.5



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

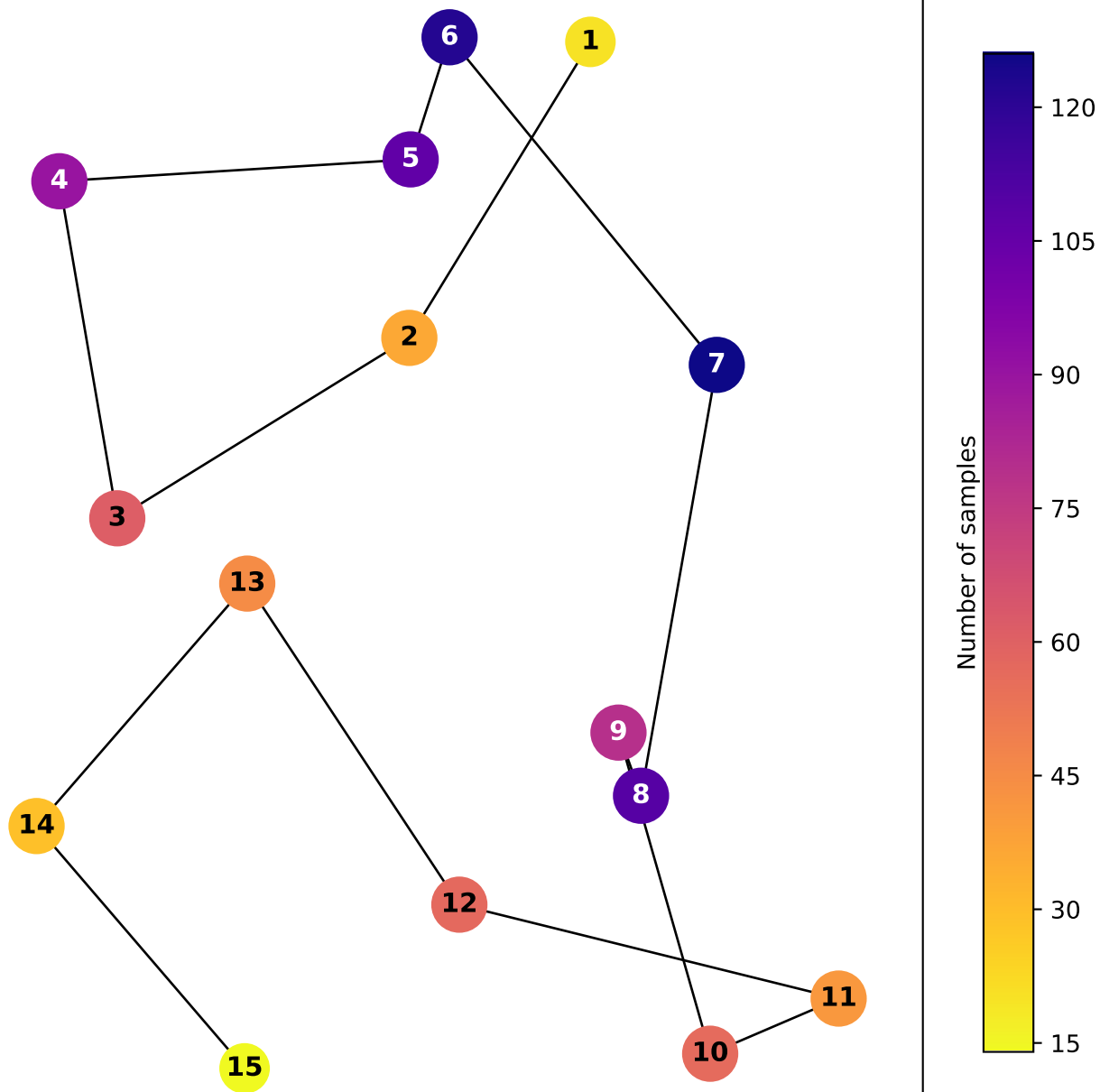
Intervals: 15 - Overlap: 0.1



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

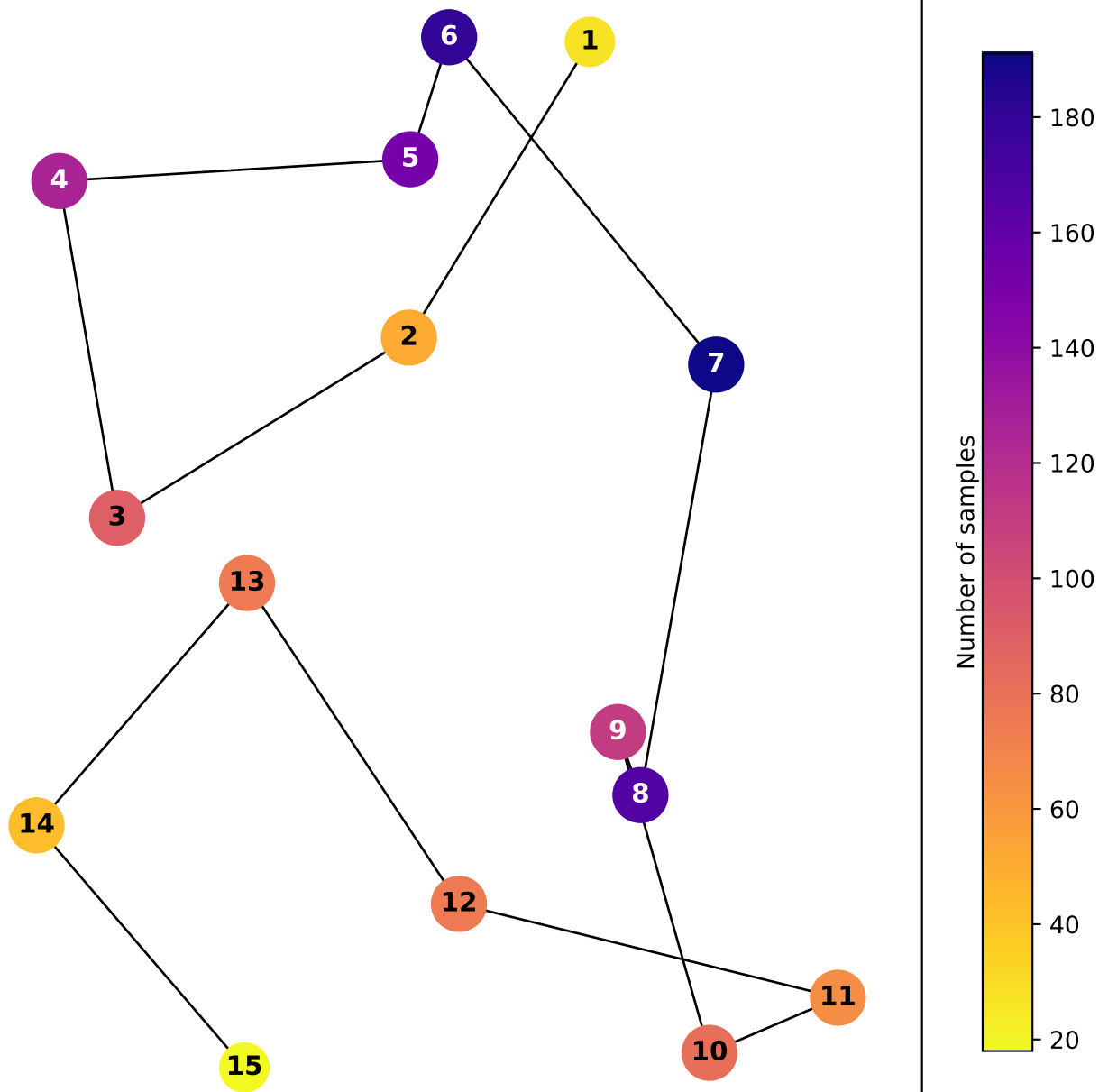
Intervals: 15 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

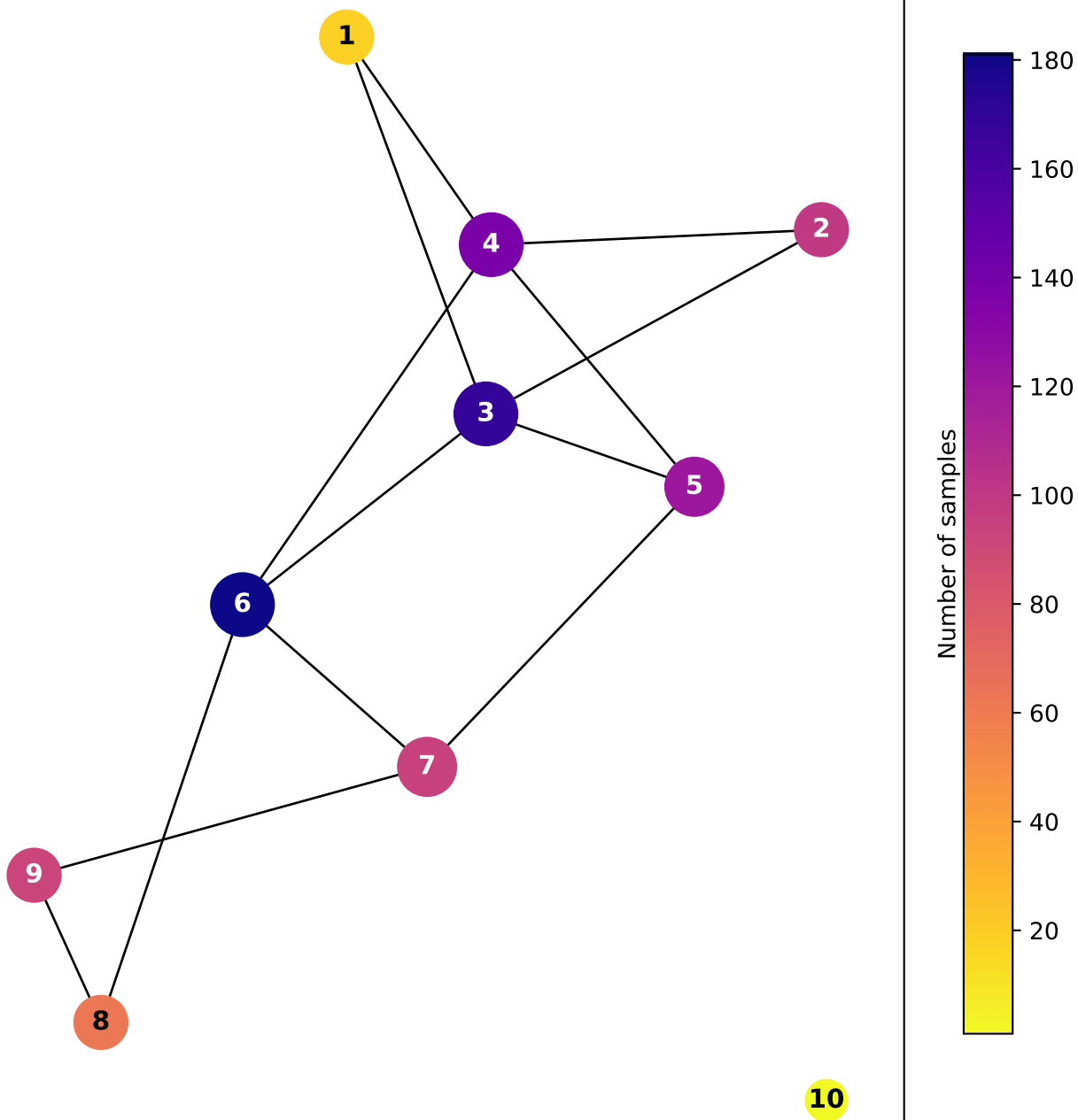
Intervals: 15 - Overlap: 0.5



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=2)

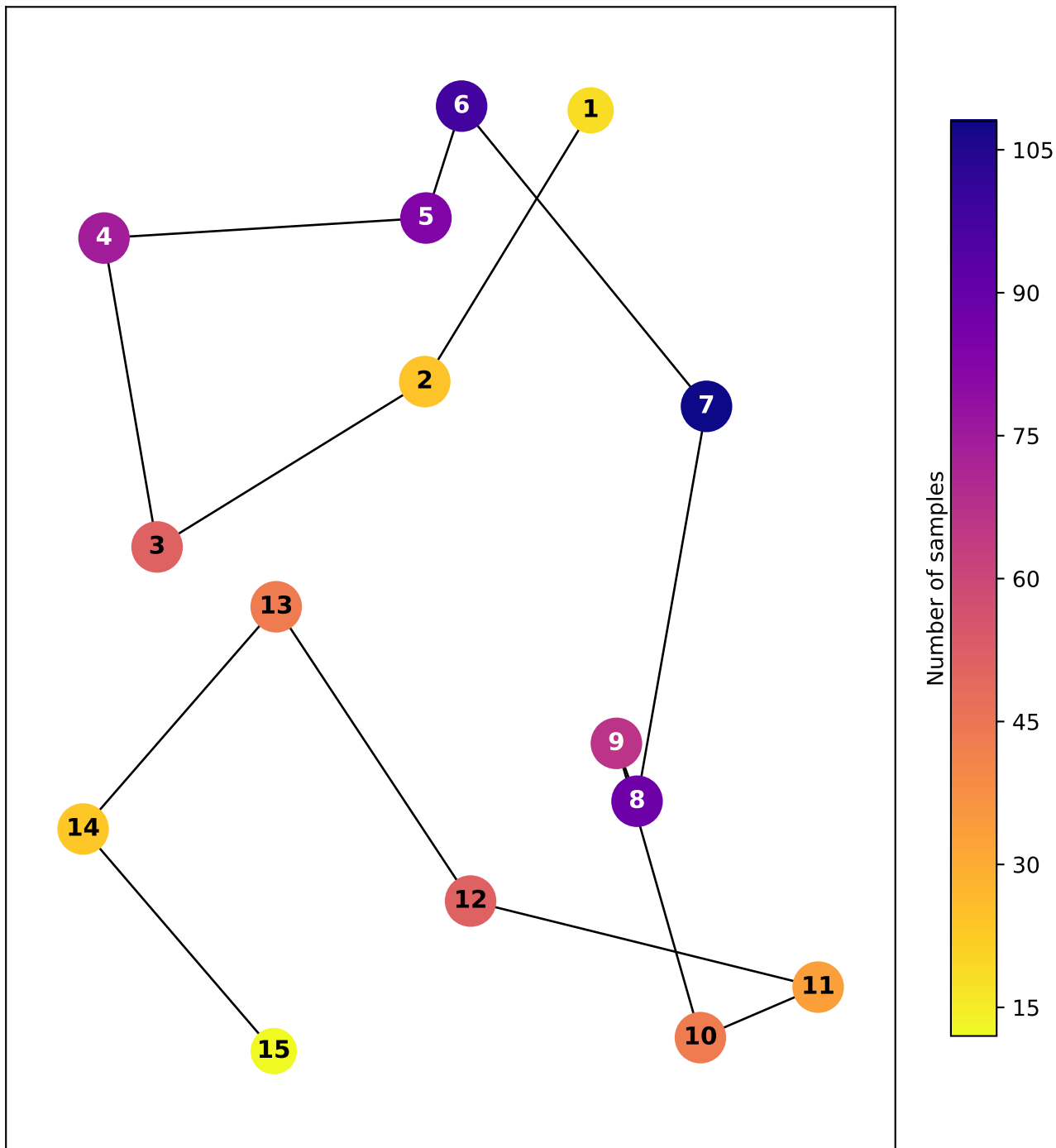
Intervals: 5 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

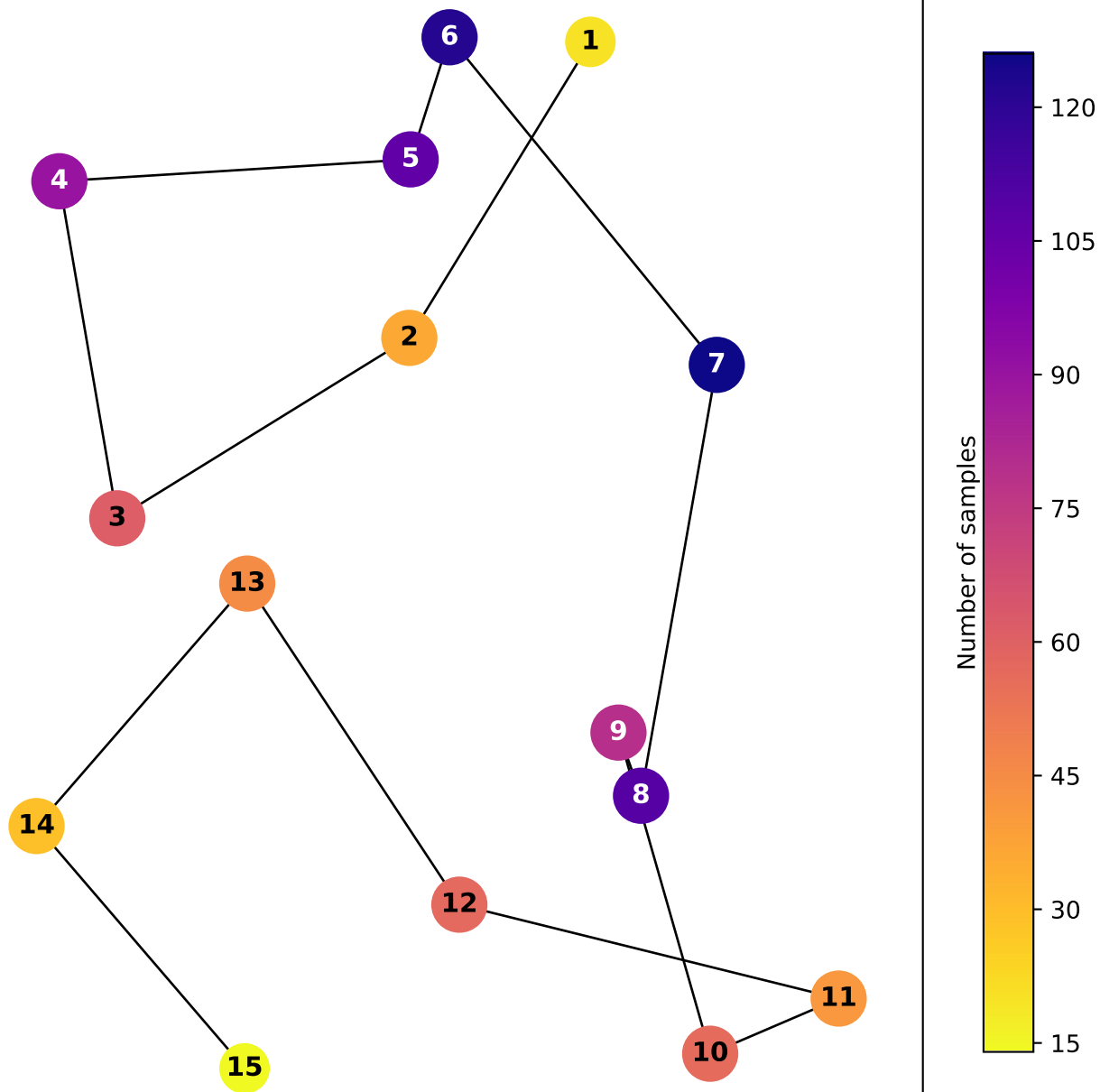
Intervals: 15 - Overlap: 0.1



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

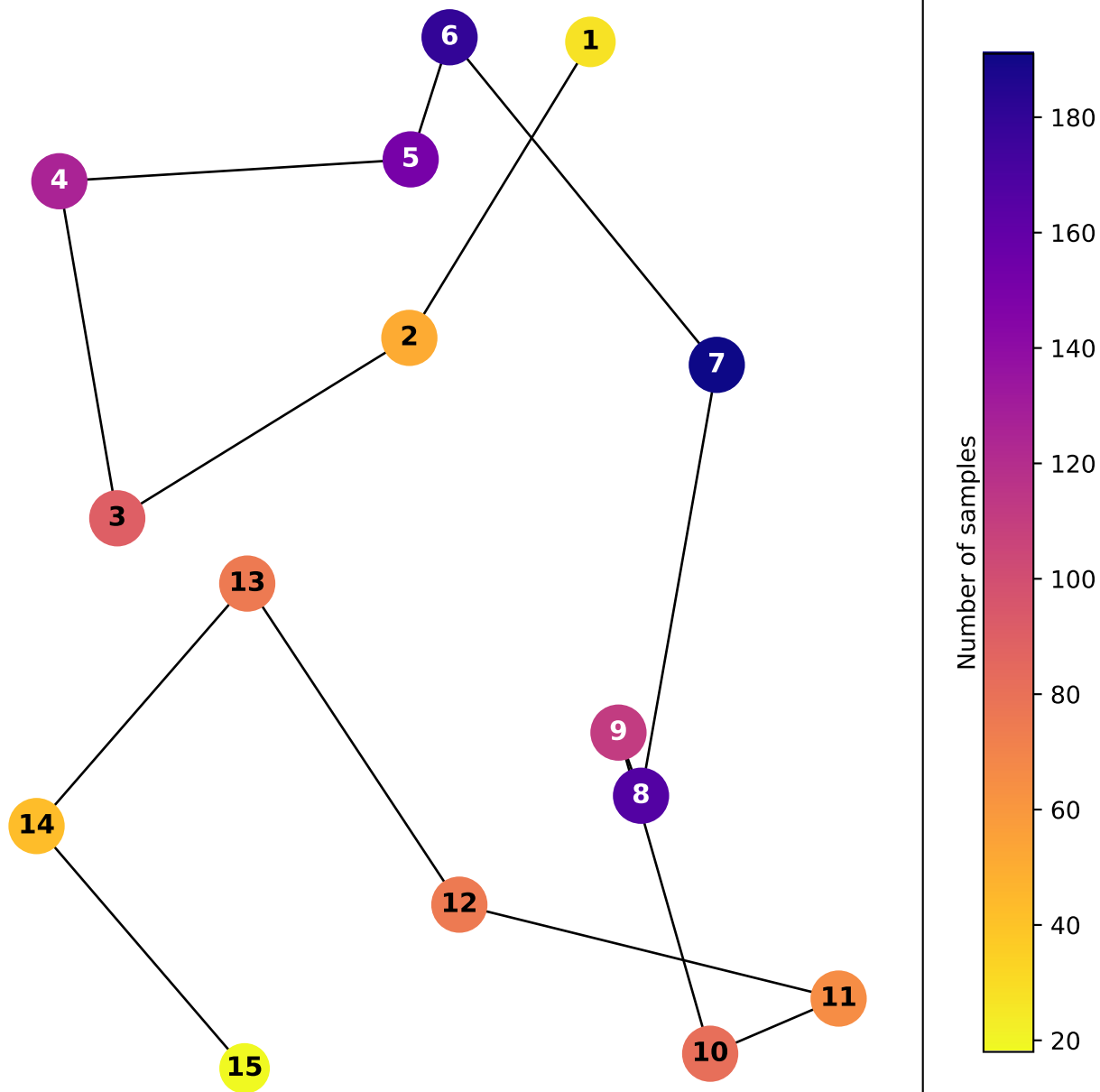
Intervals: 15 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

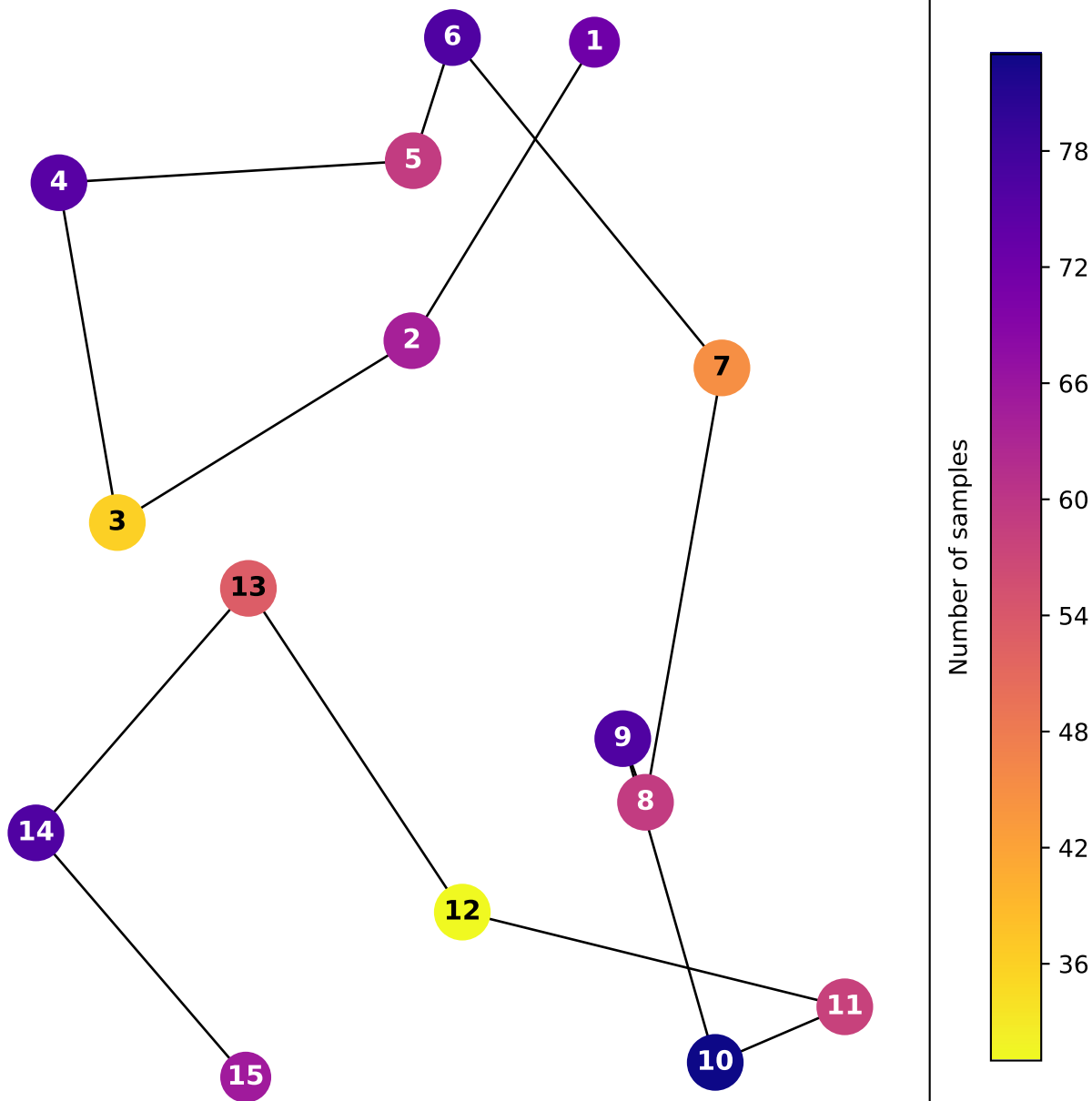
Intervals: 15 - Overlap: 0.5



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

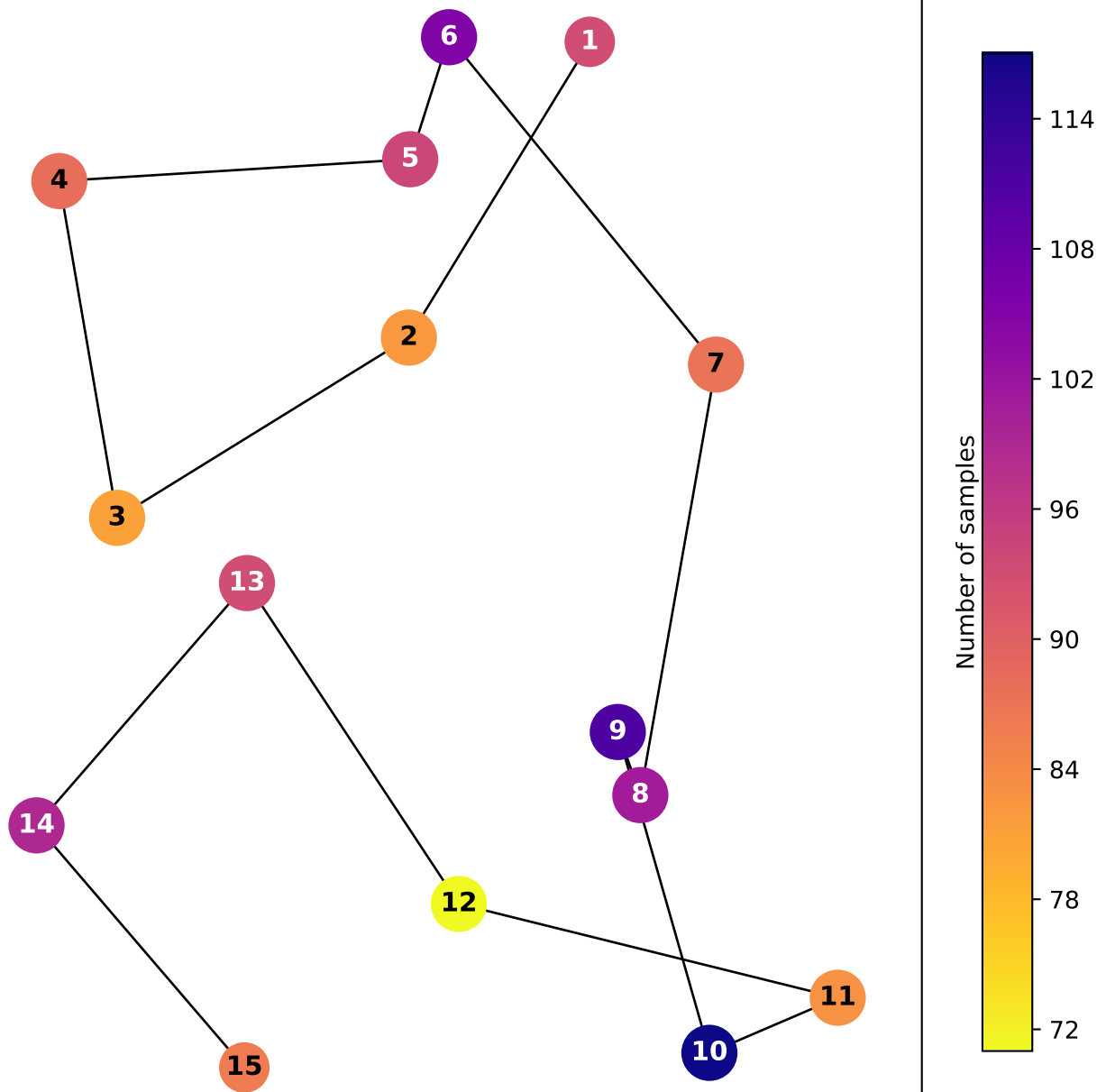
Intervals: 15 - Overlap: 0.25



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

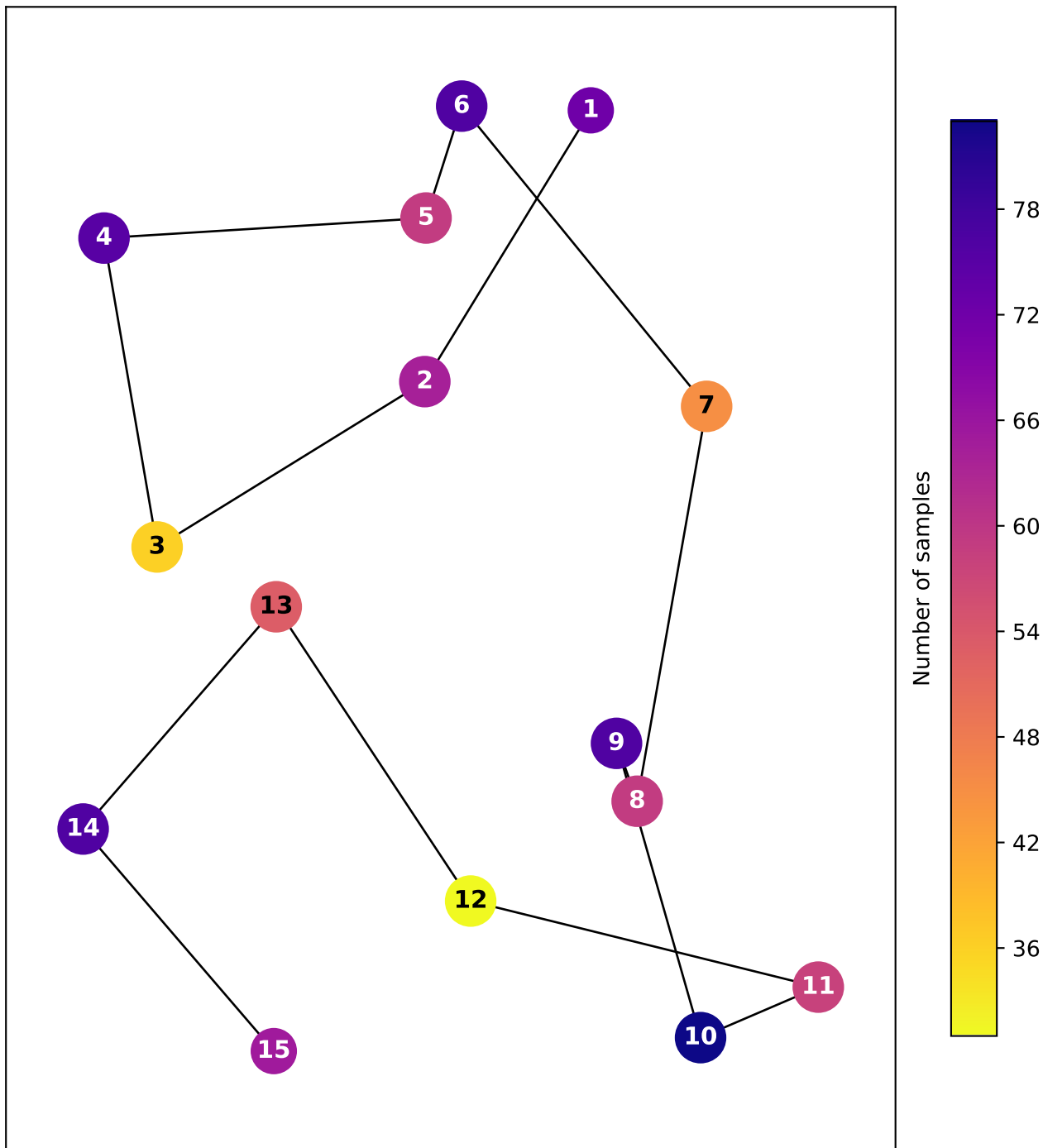
Intervals: 15 - Overlap: 0.5



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

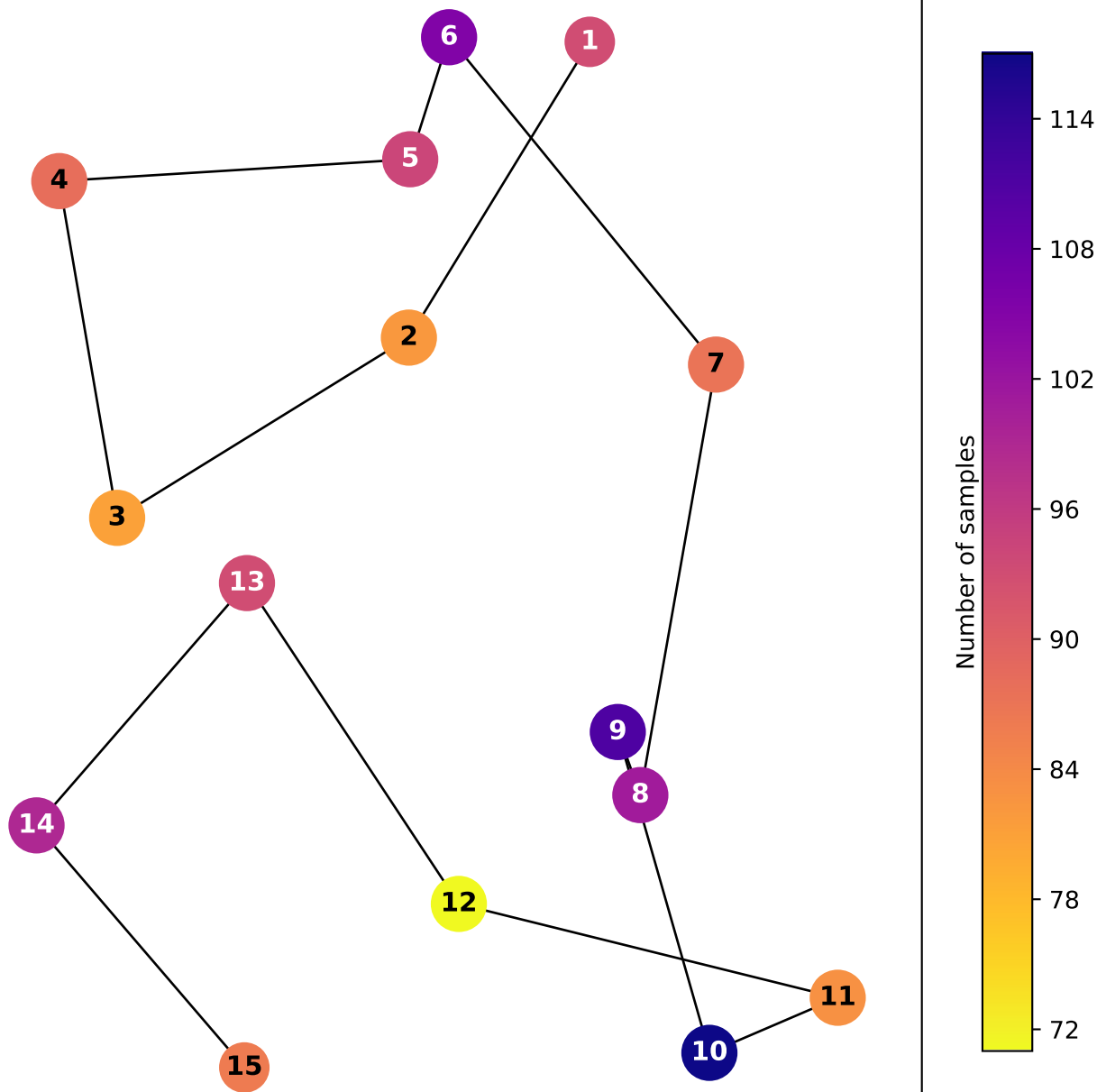
Intervals: 15 - Overlap: 0.25



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

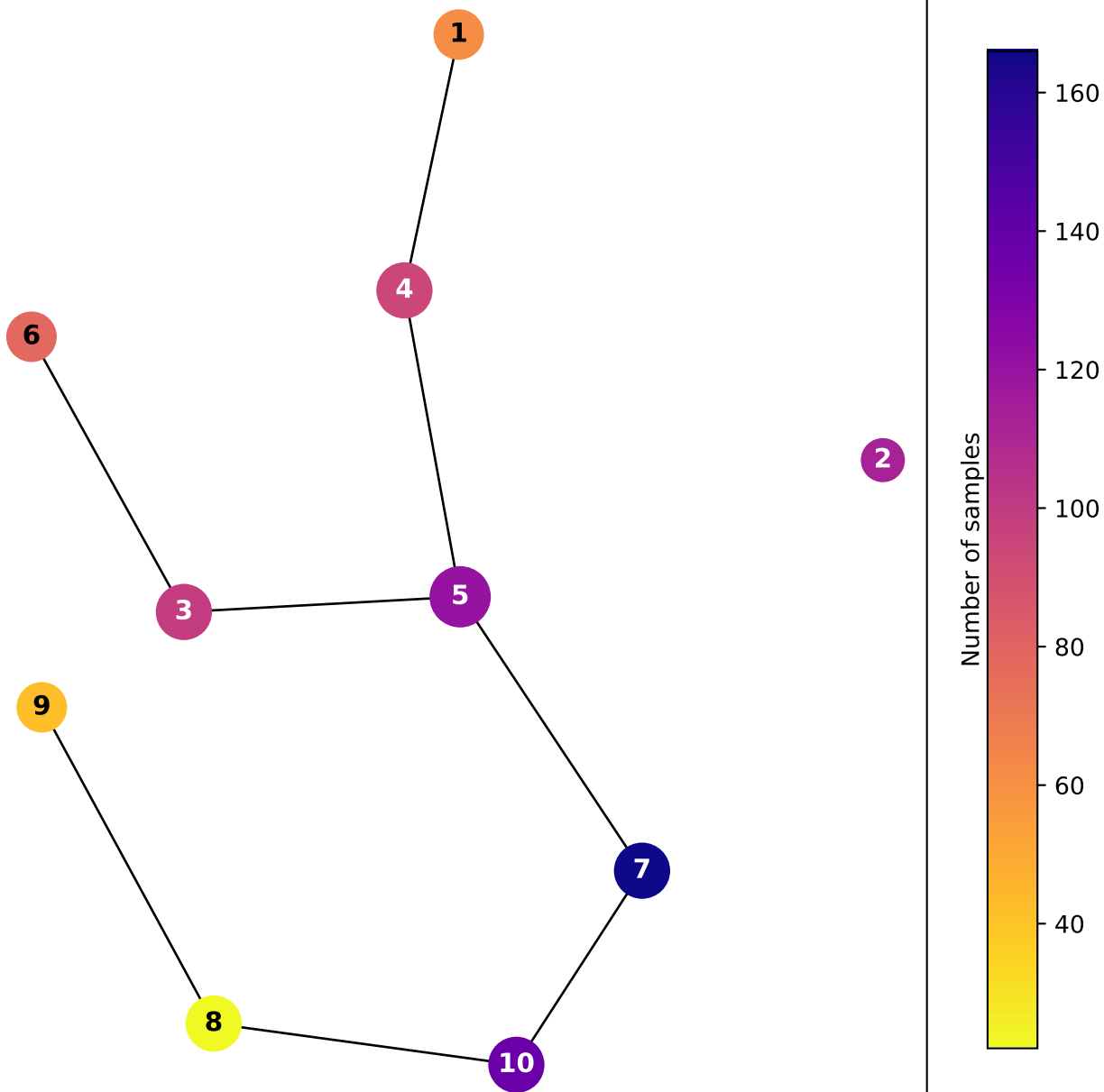
Intervals: 15 - Overlap: 0.5



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=2)

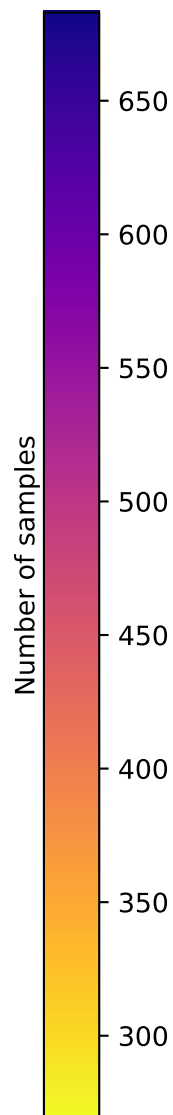
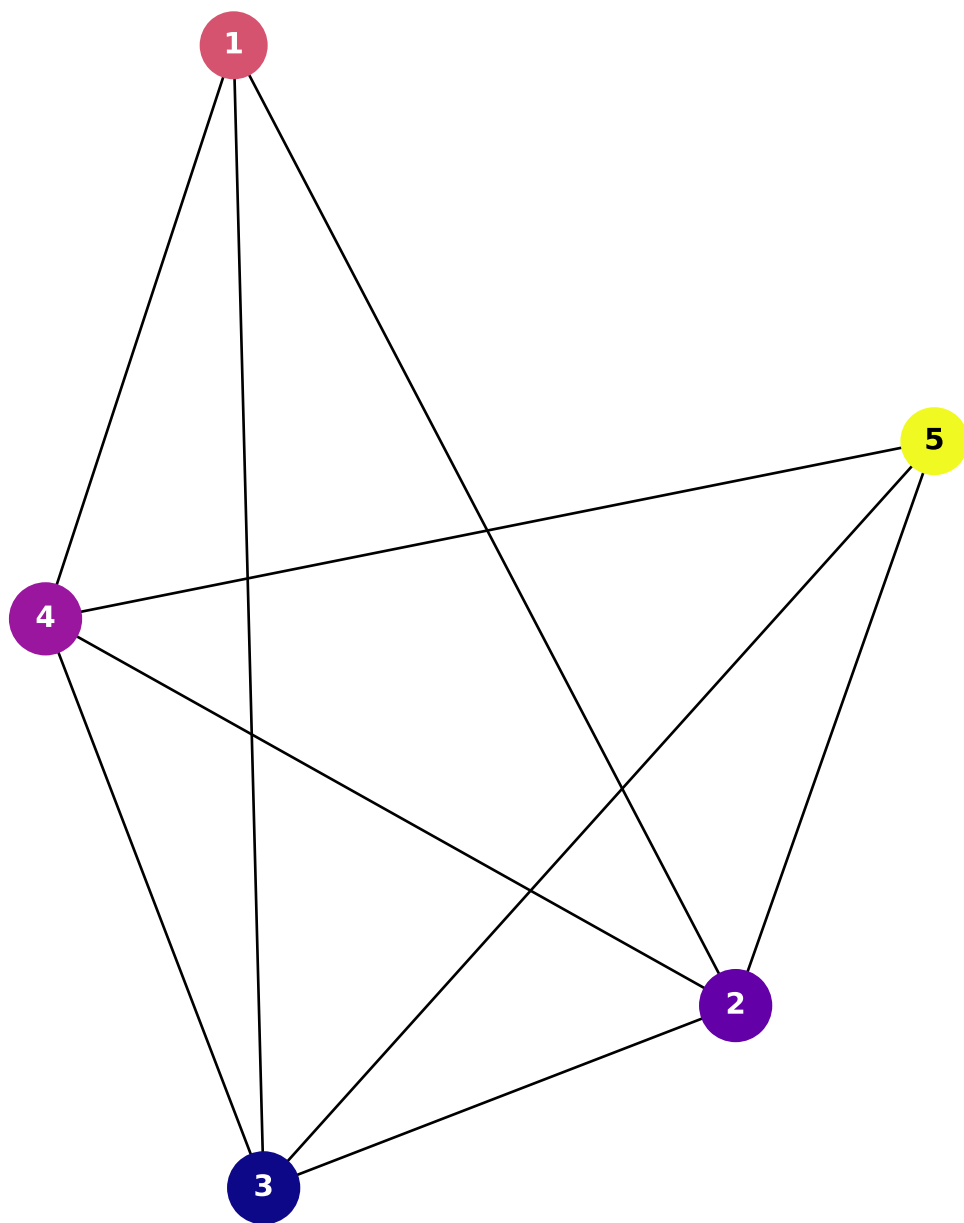
Intervals: 5 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

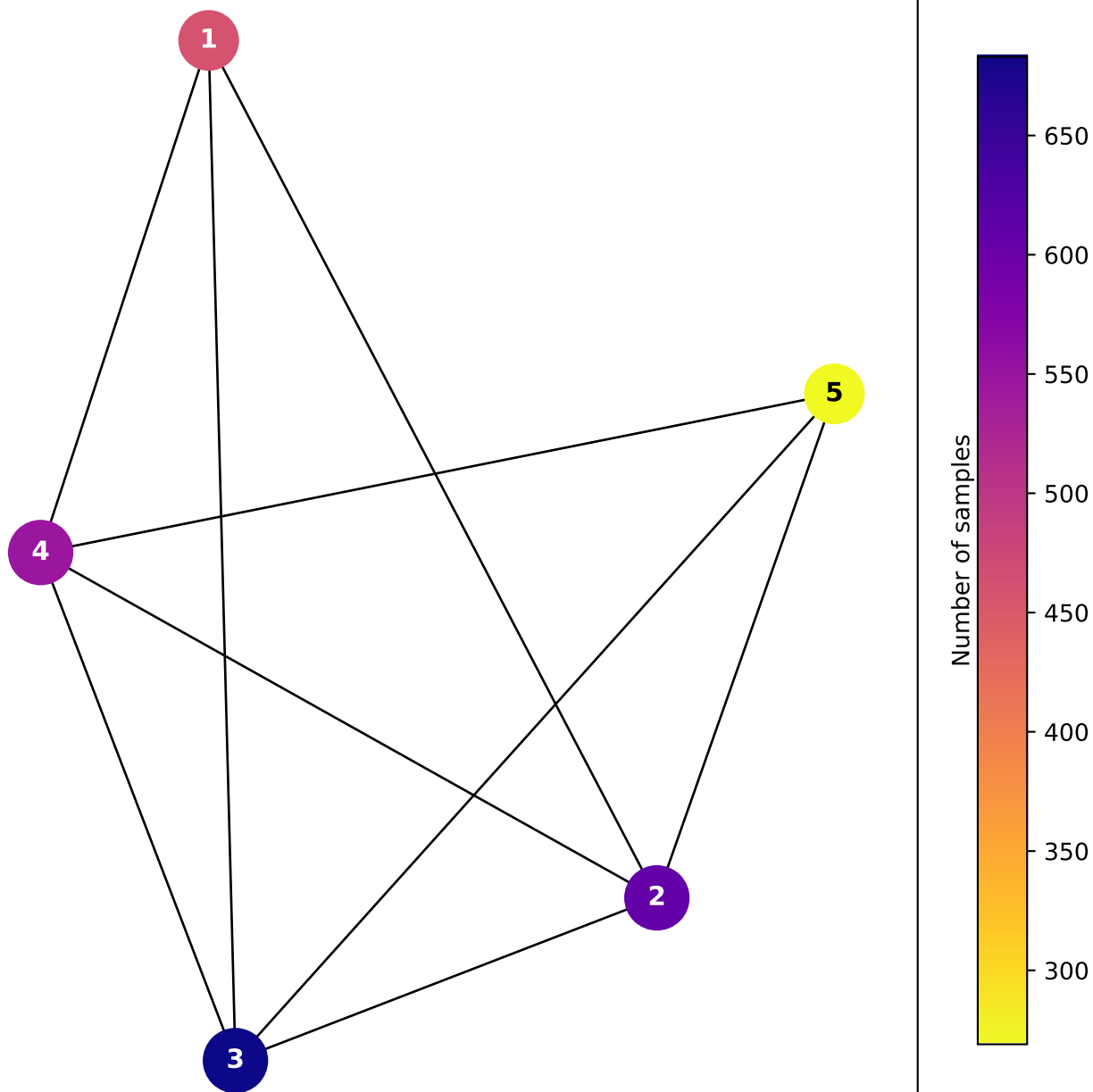
Intervals: 5 - Overlap: 0.75



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

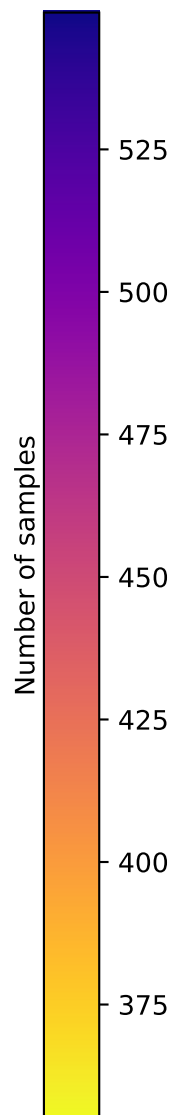
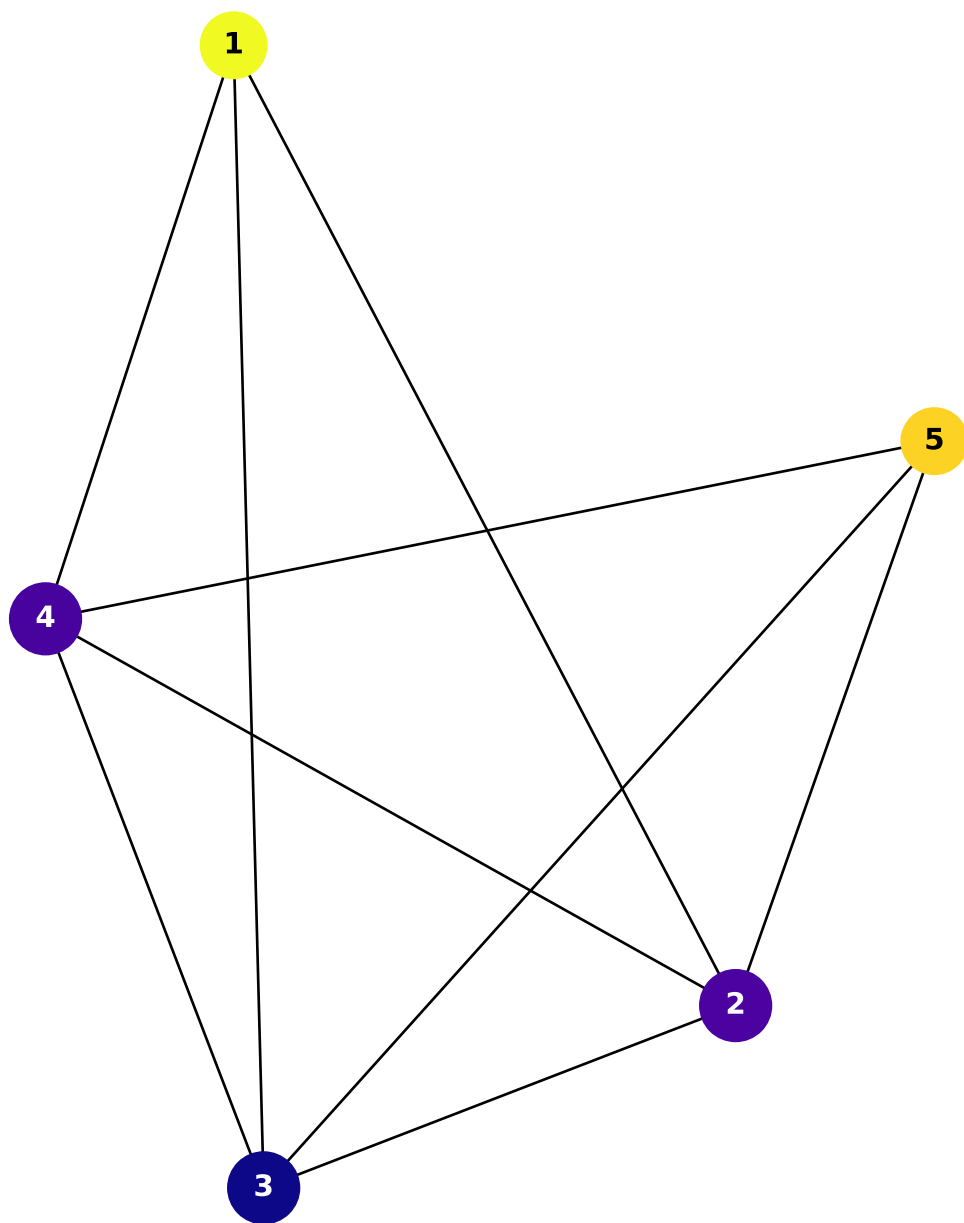
Intervals: 5 - Overlap: 0.75



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

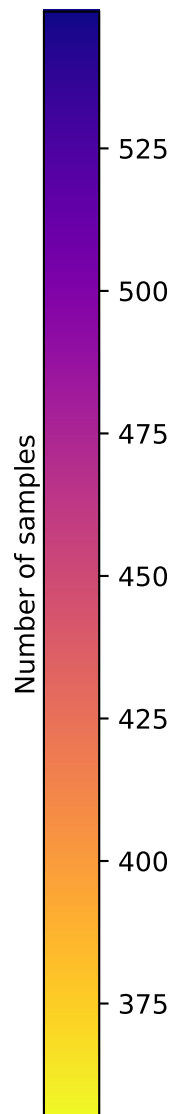
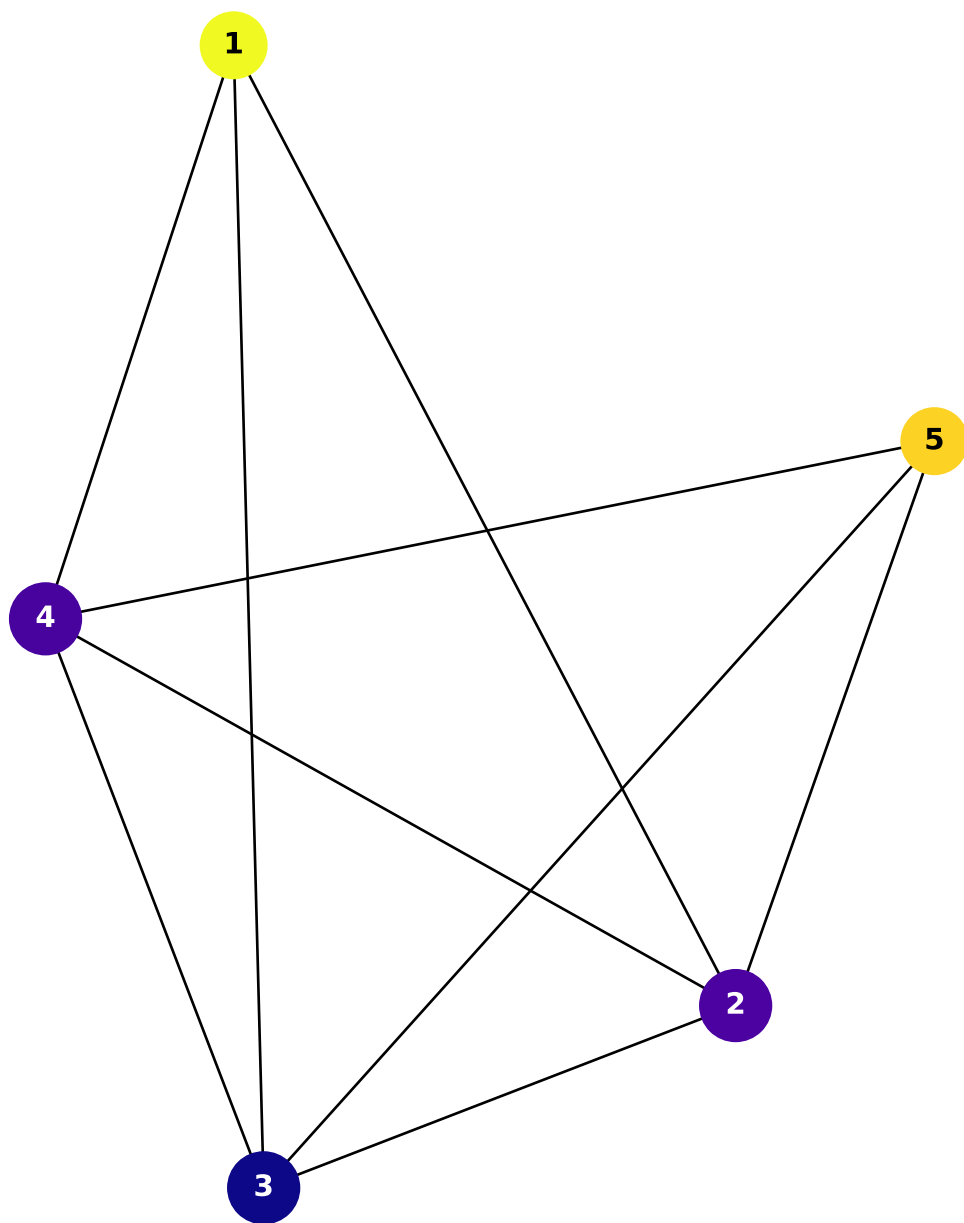
Intervals: 5 - Overlap: 0.75



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

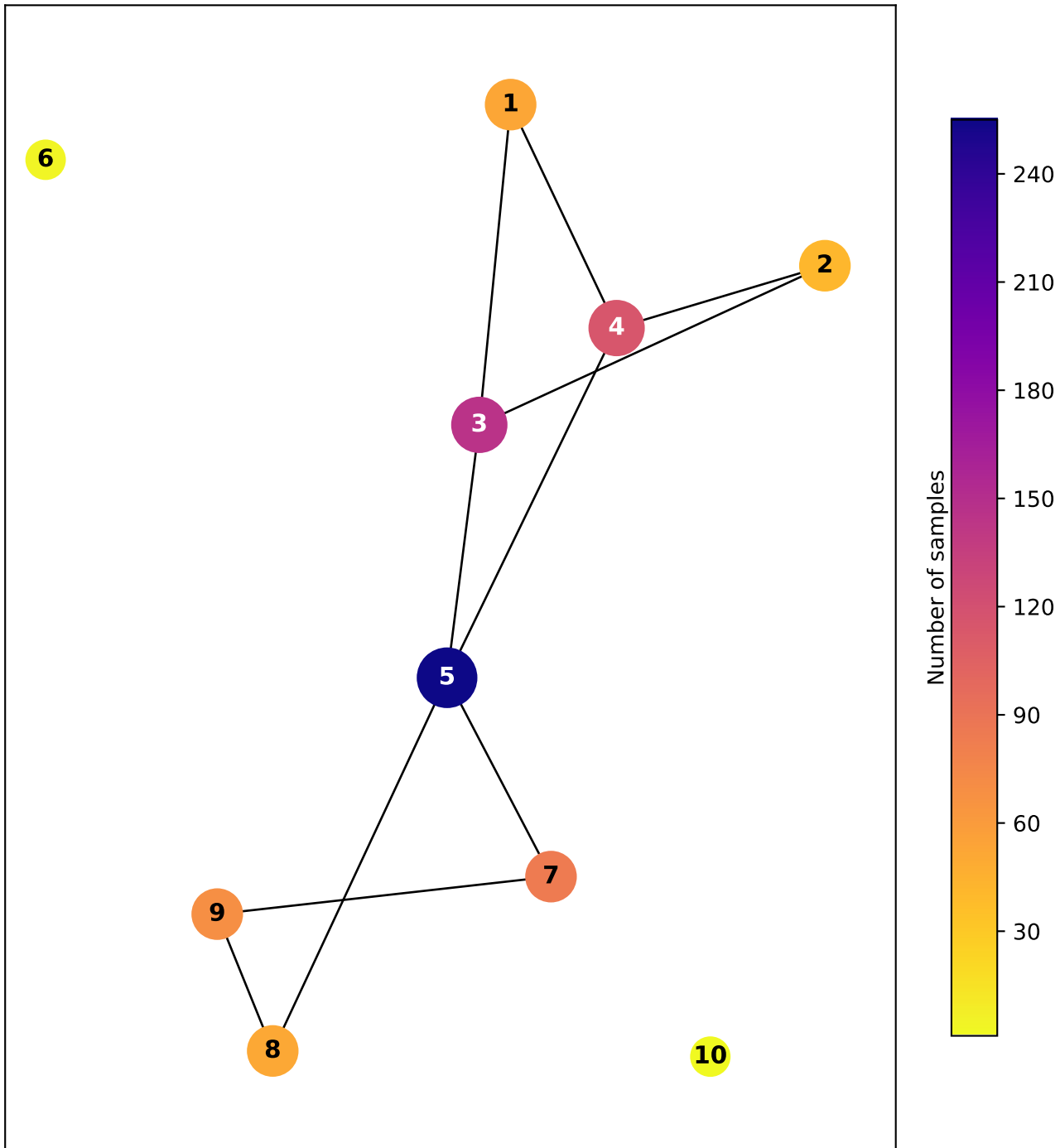
Intervals: 5 - Overlap: 0.75



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=2)

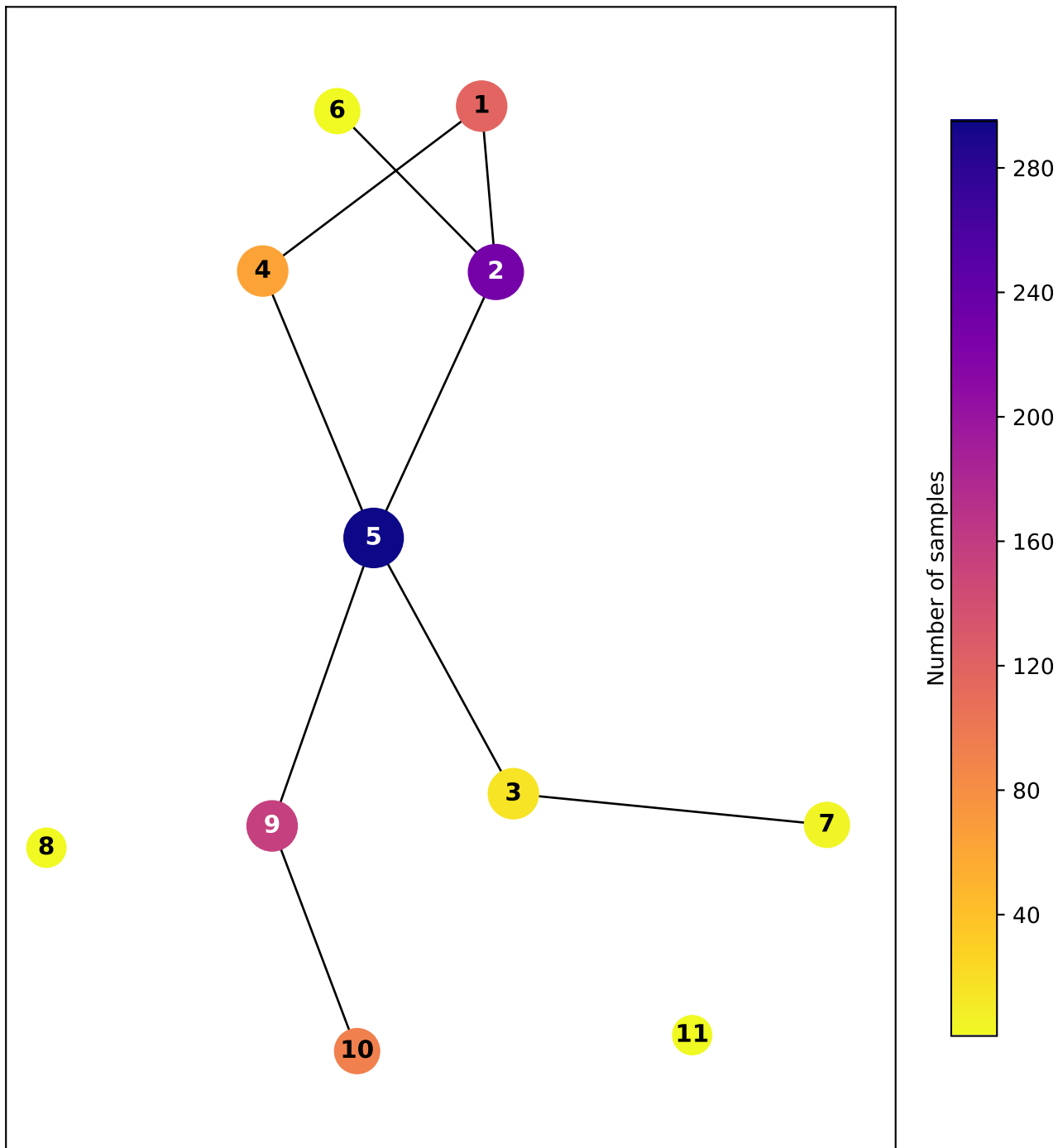
Intervals: 5 - Overlap: 0.1



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=30.0)

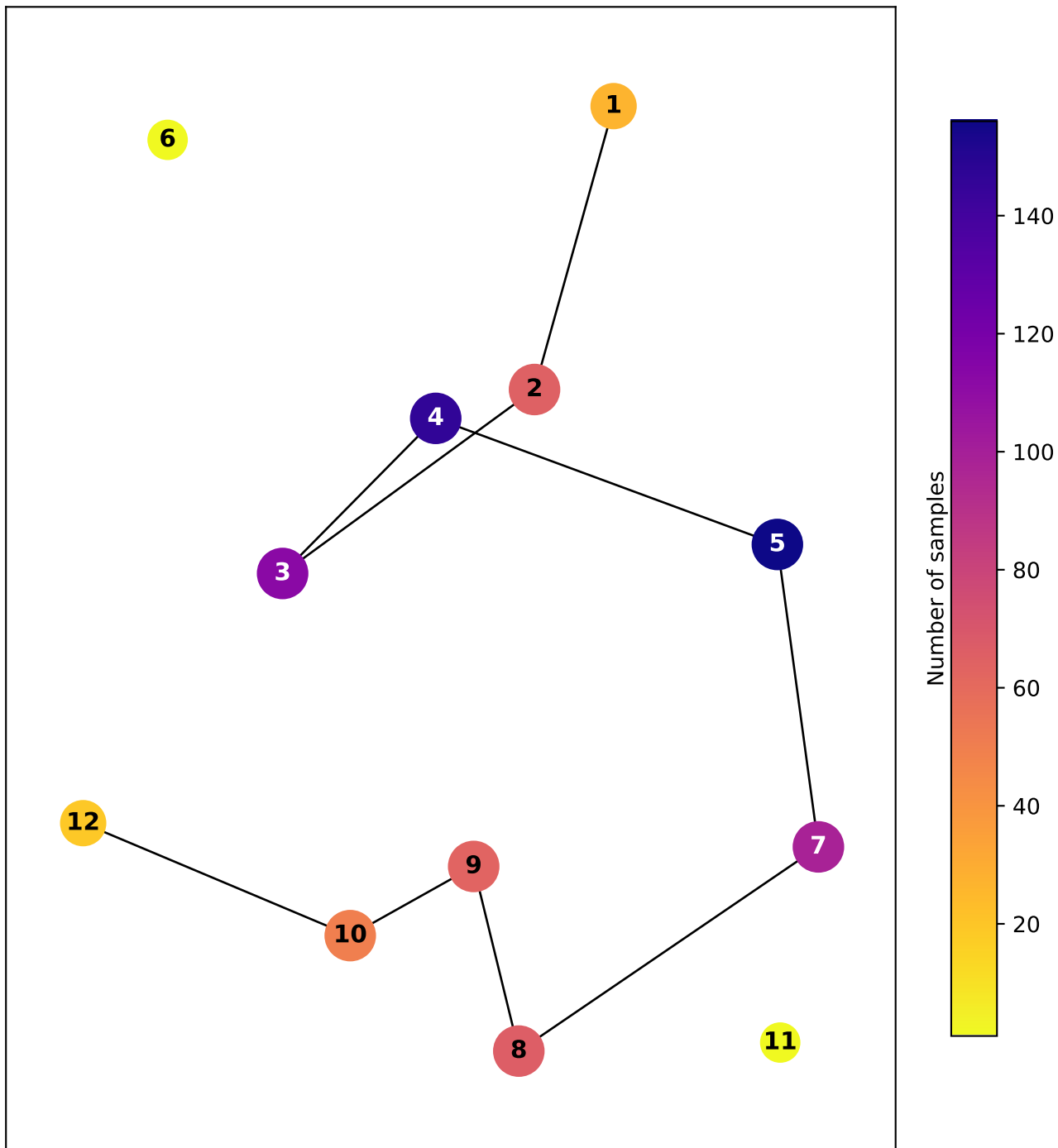
Intervals: 5 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=30.0)

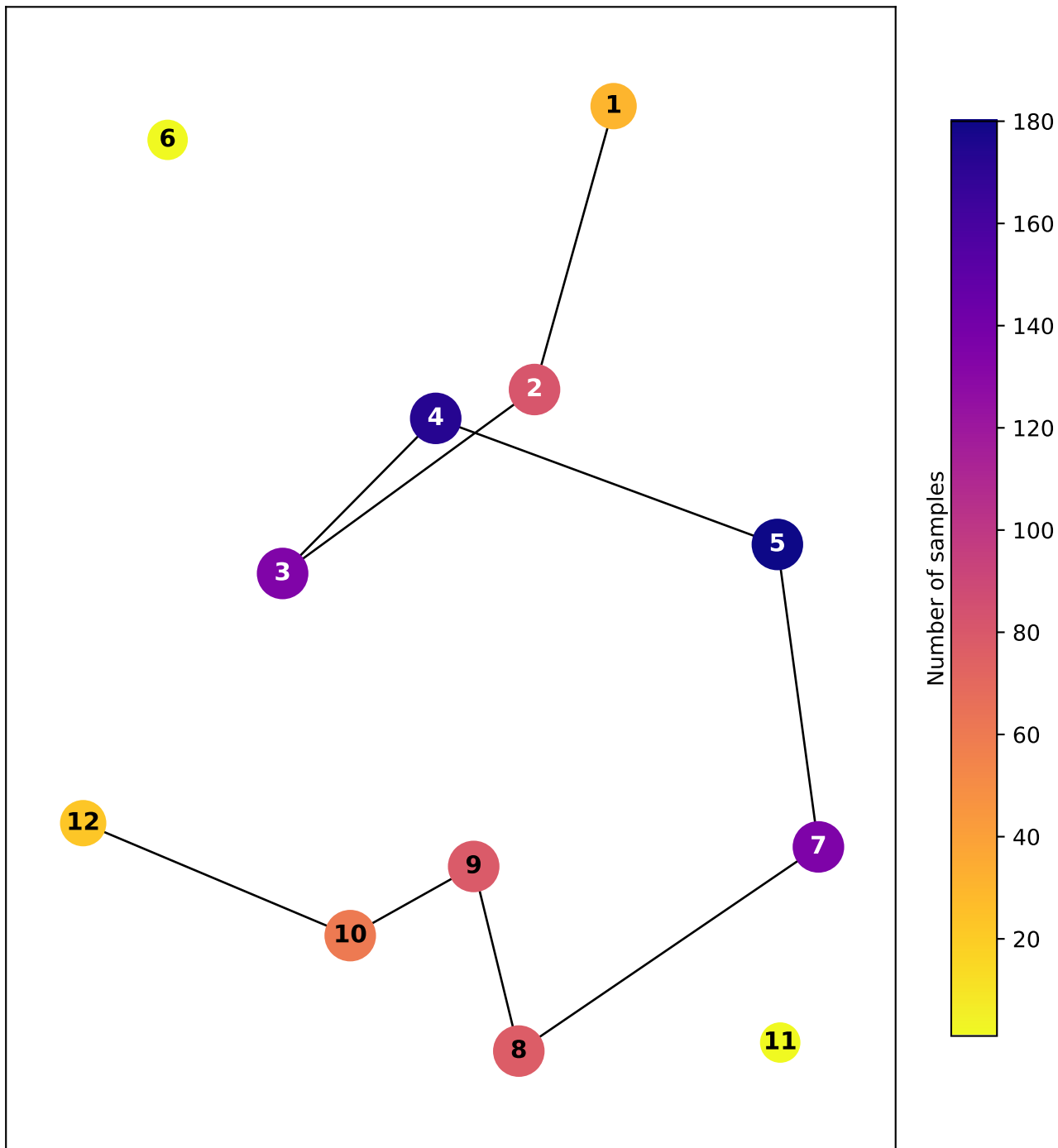
Intervals: 10 - Overlap: 0.1



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=30.0)

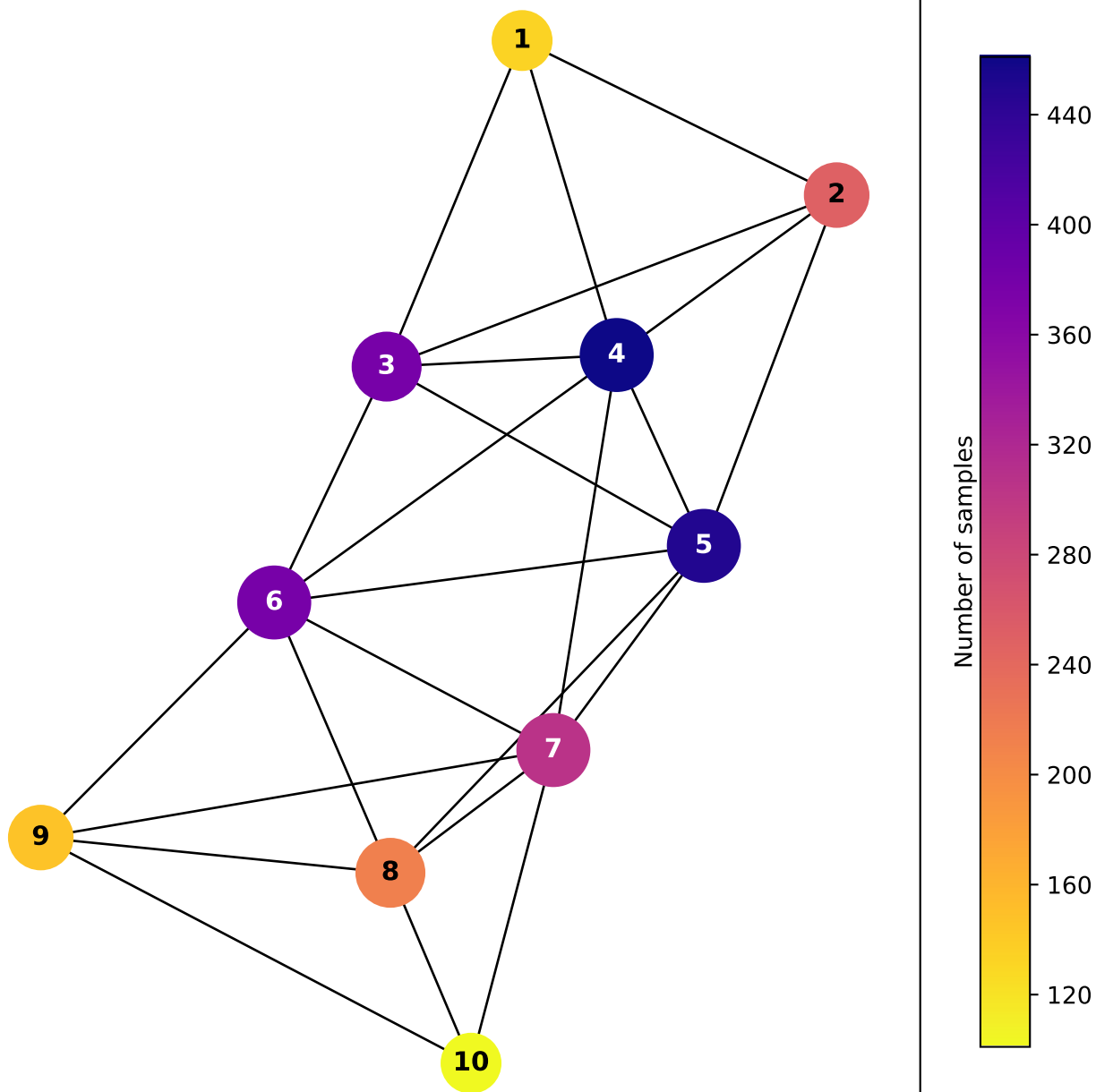
Intervals: 10 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

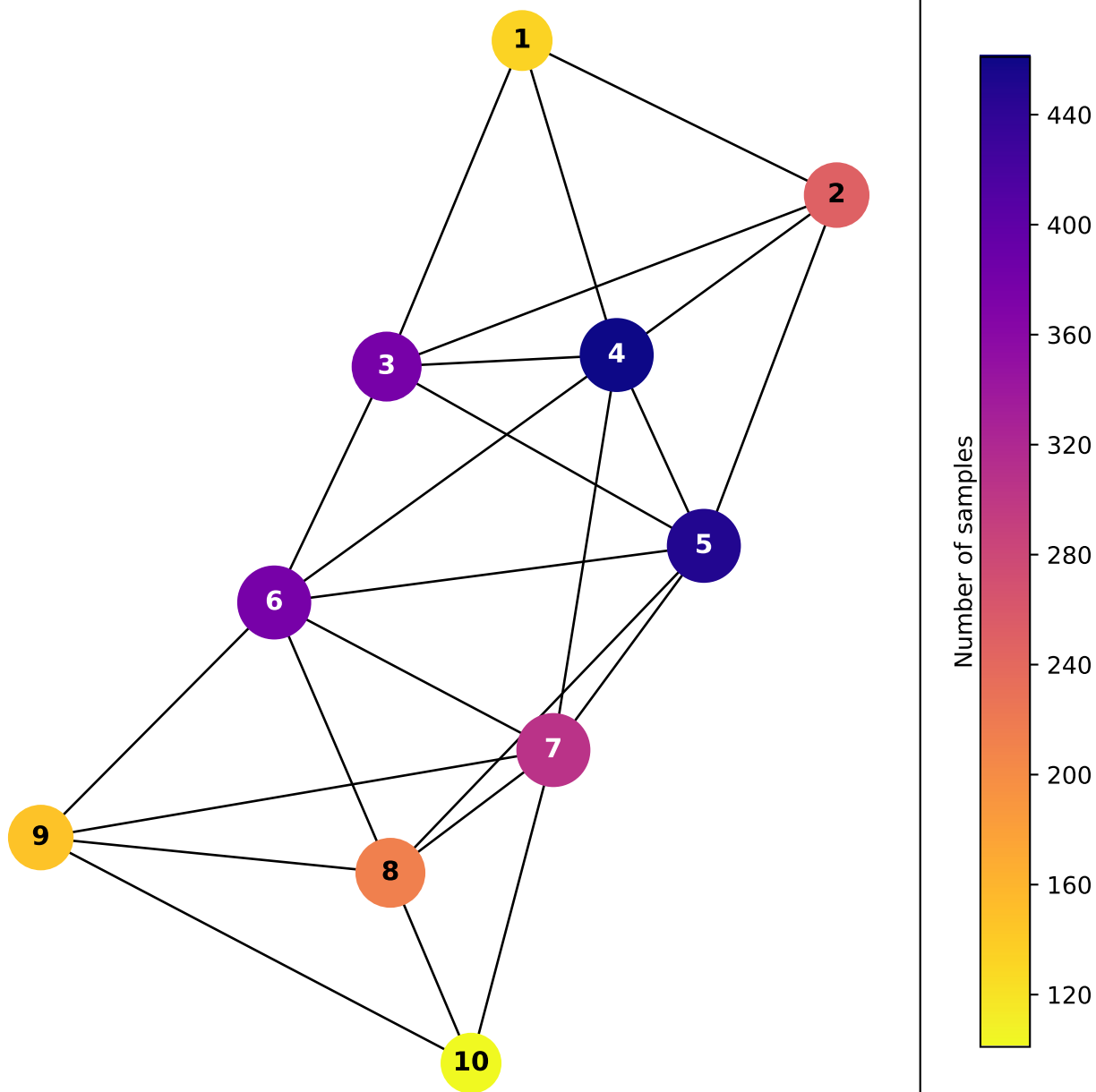
Intervals: 10 - Overlap: 0.75



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

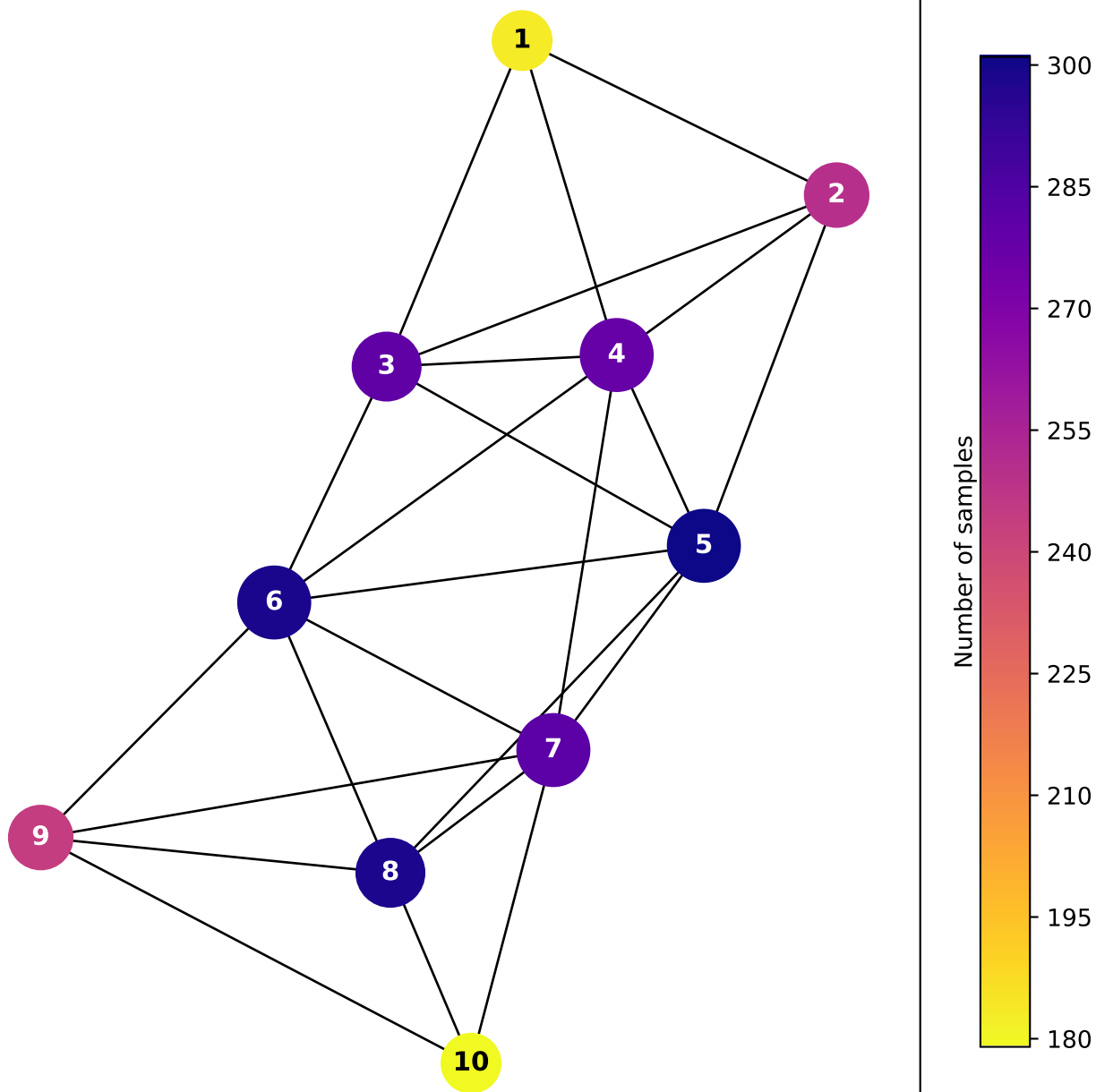
Intervals: 10 - Overlap: 0.75



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

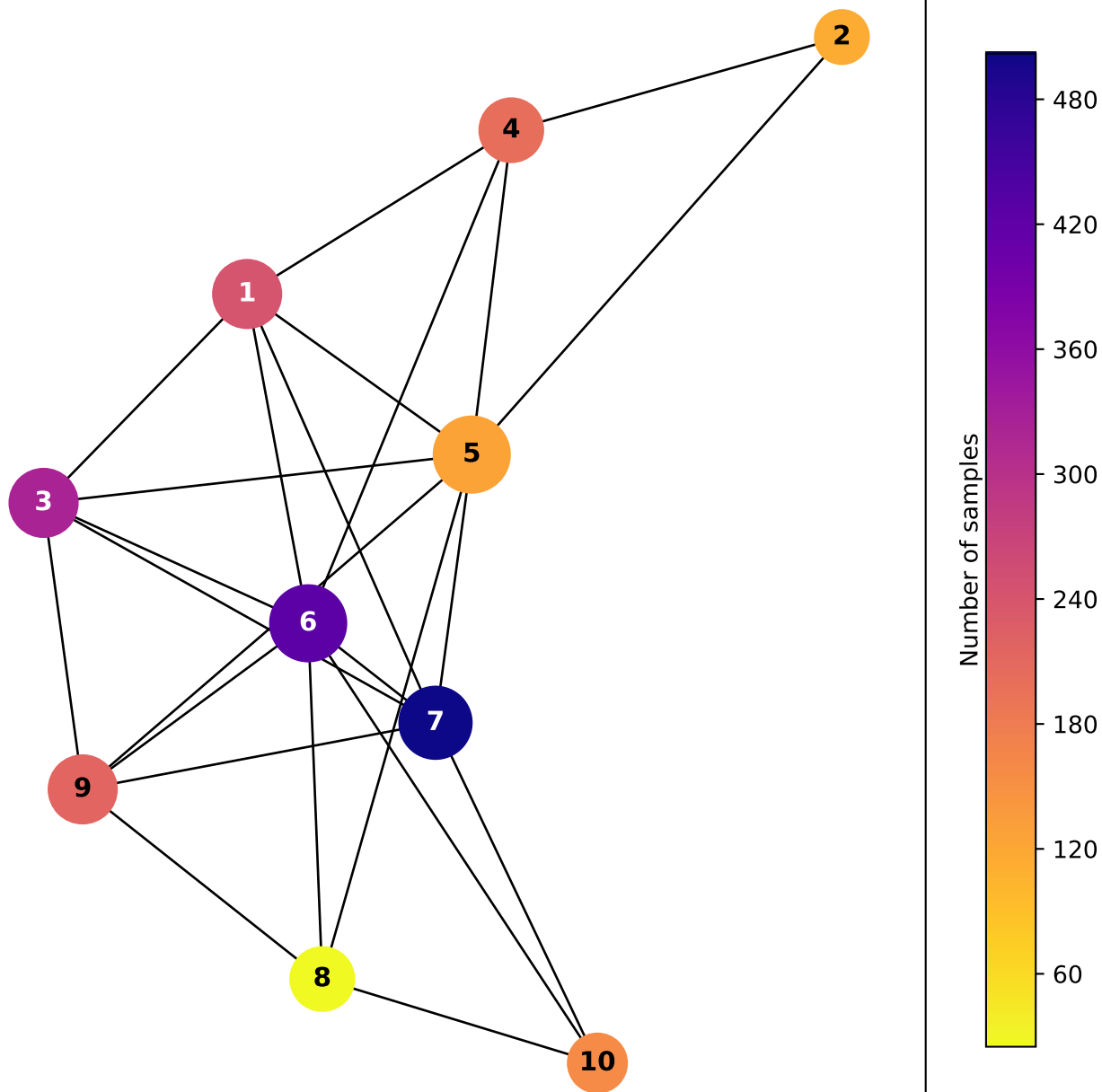
Intervals: 10 - Overlap: 0.75



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=2)

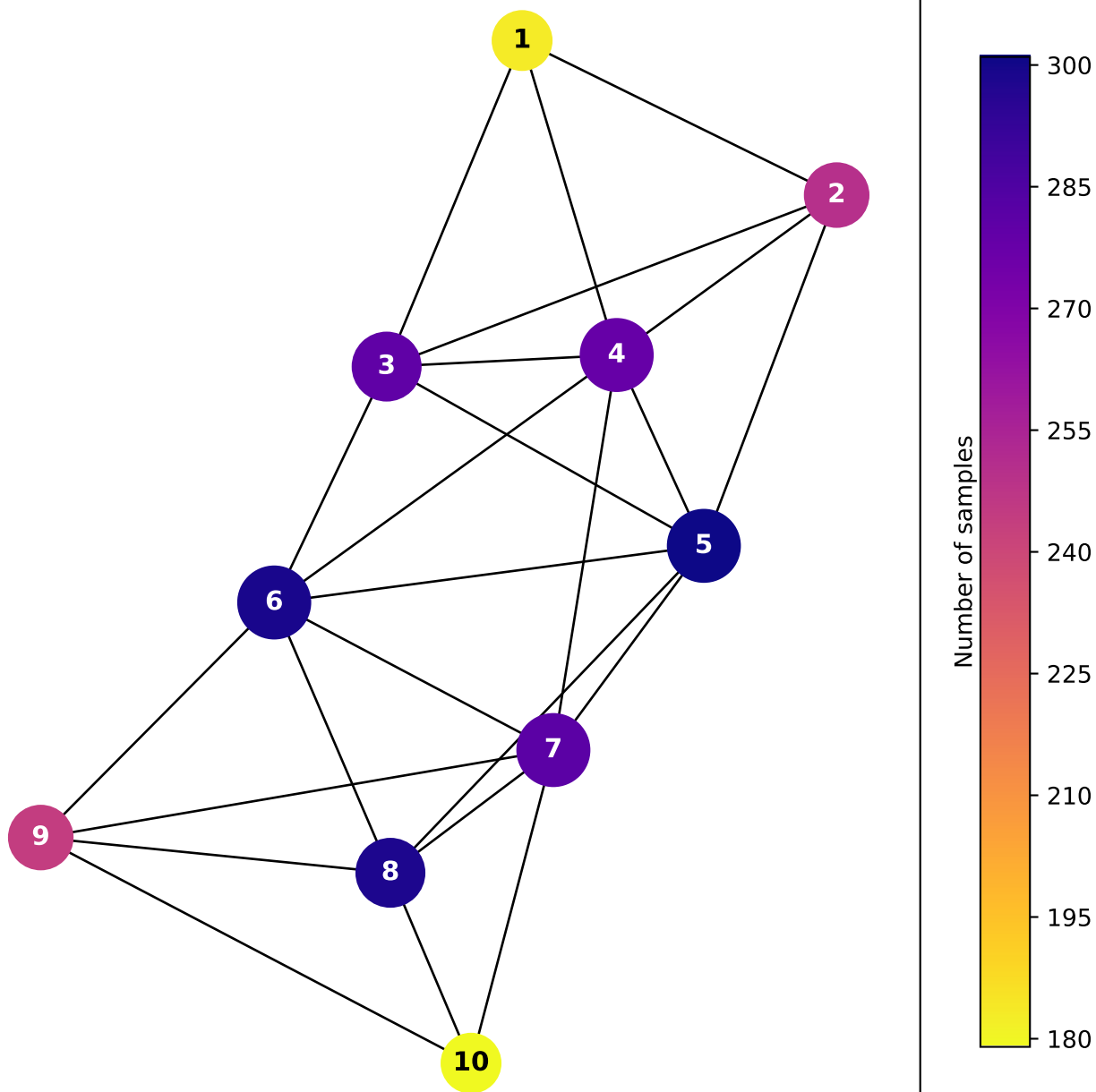
Intervals: 5 - Overlap: 0.75



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

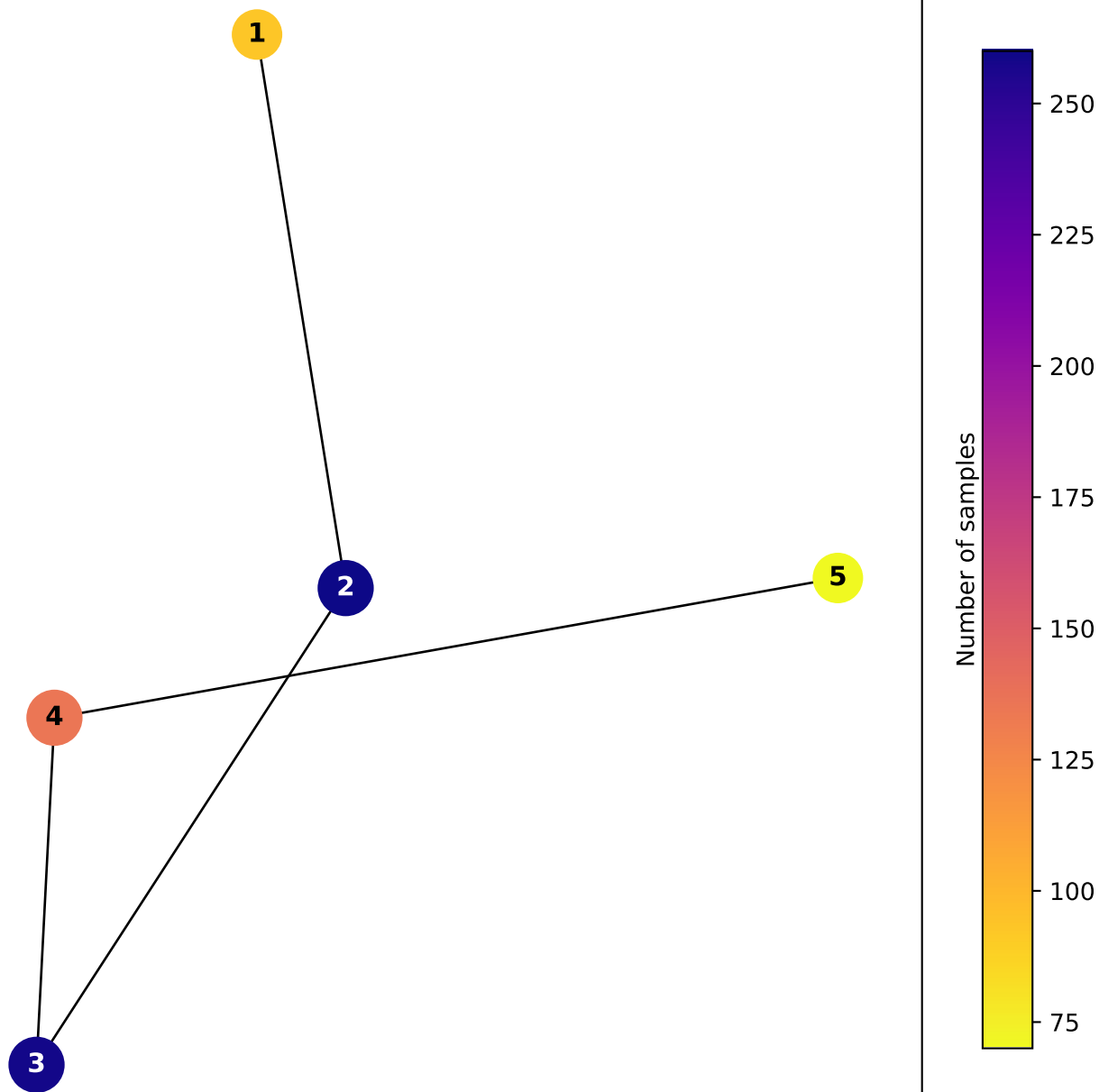
Intervals: 10 - Overlap: 0.75



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

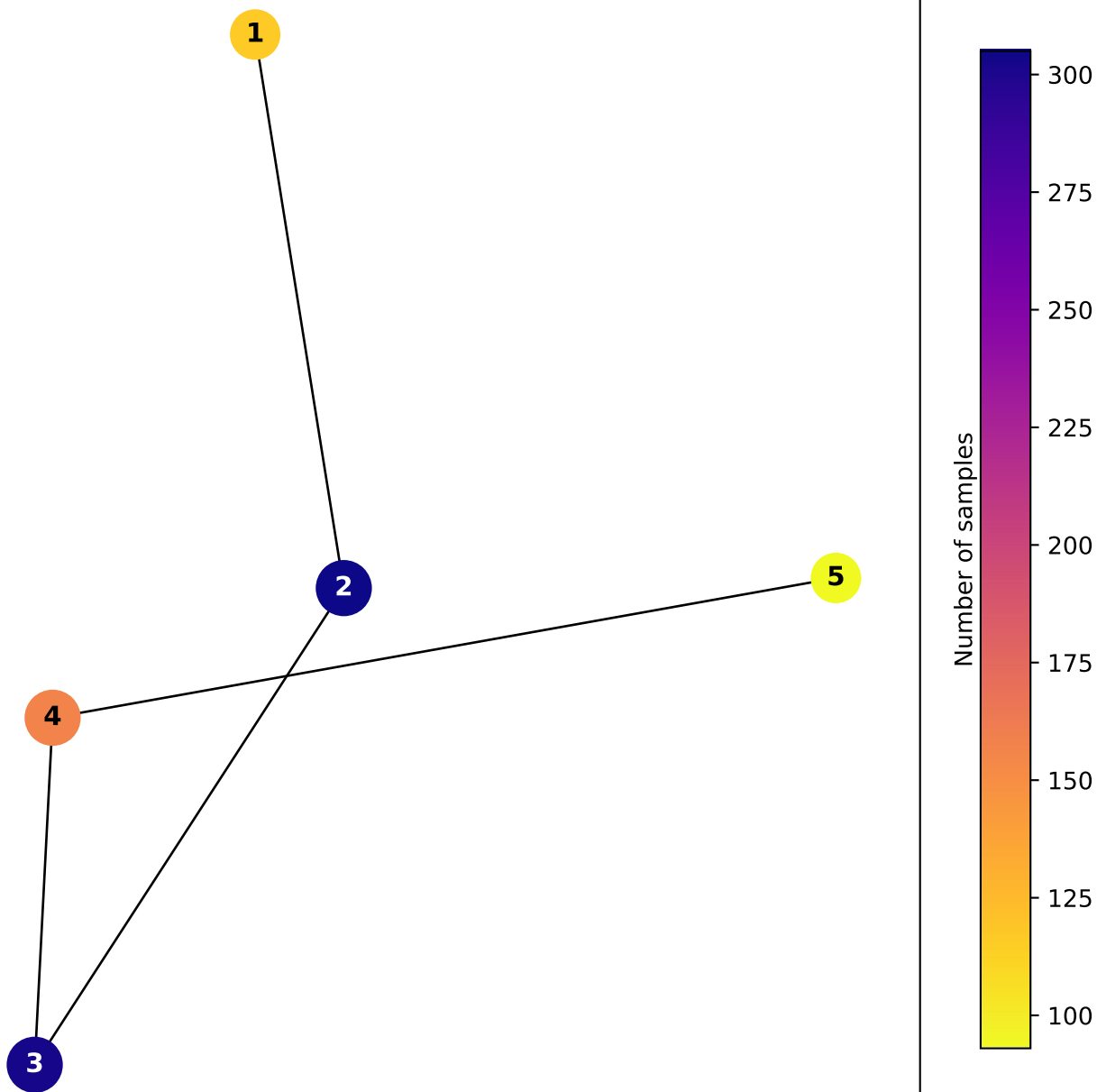
Intervals: 5 - Overlap: 0.1



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

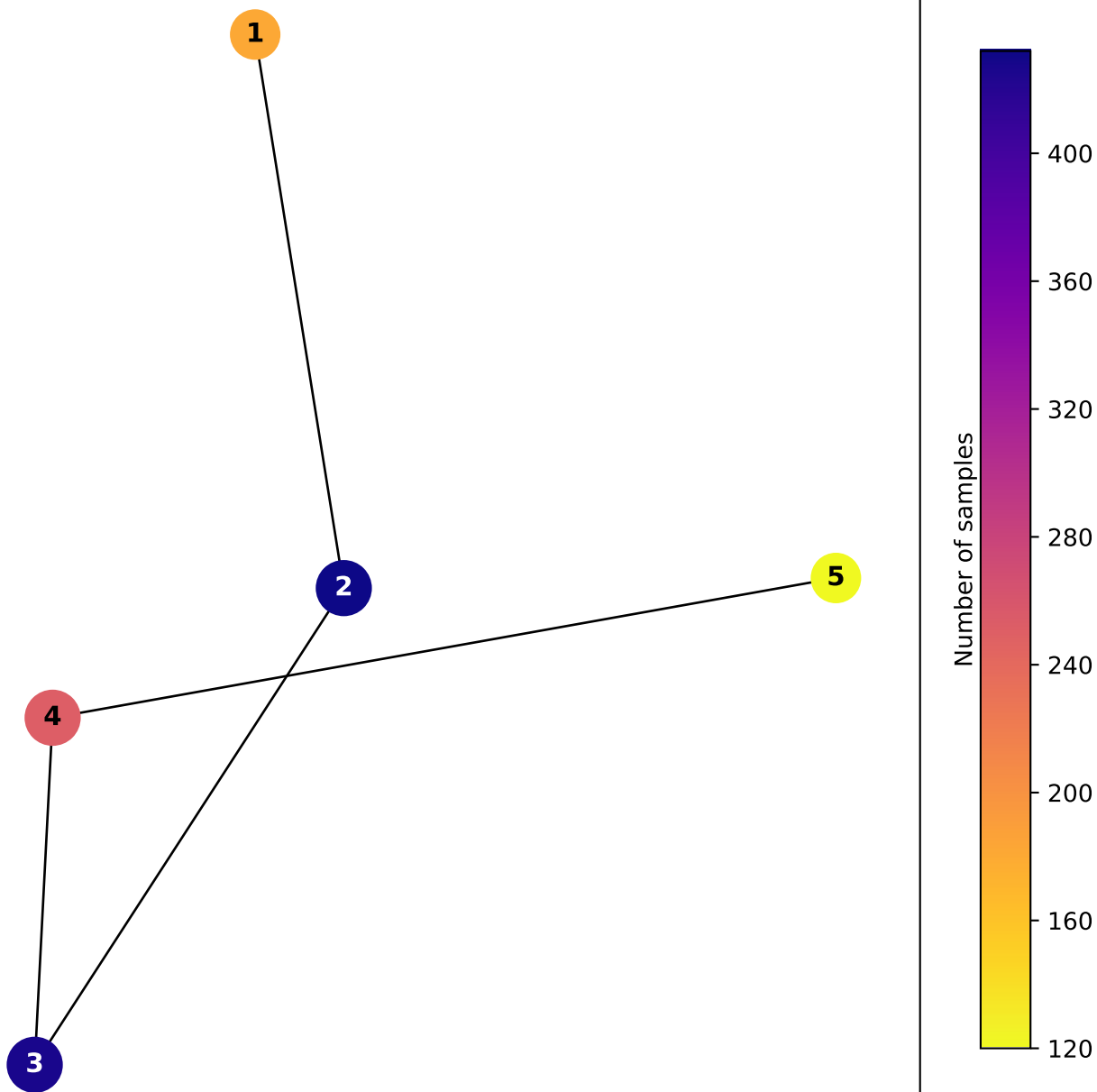
Intervals: 5 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

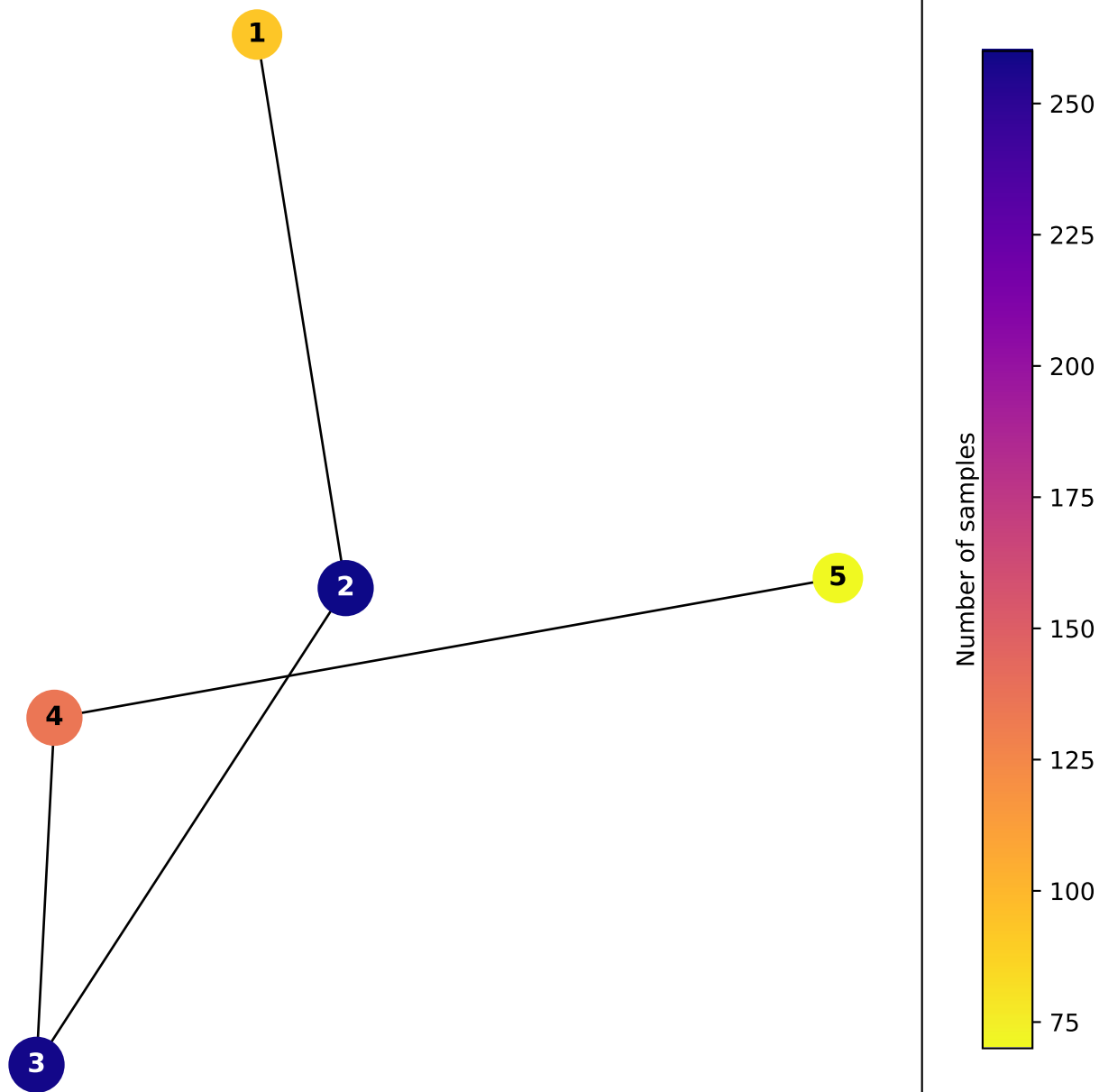
Intervals: 5 - Overlap: 0.5



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

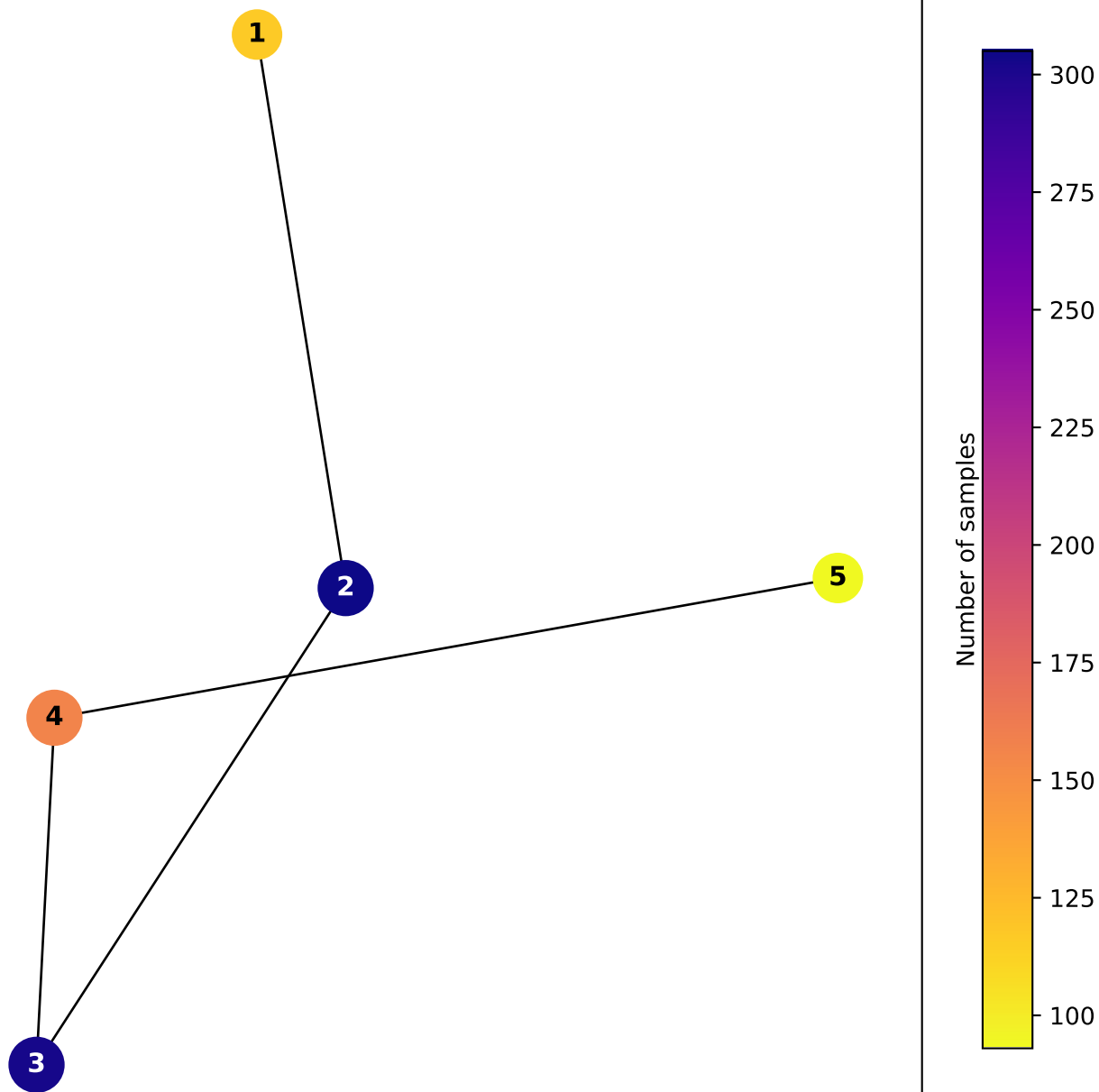
Intervals: 5 - Overlap: 0.1



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

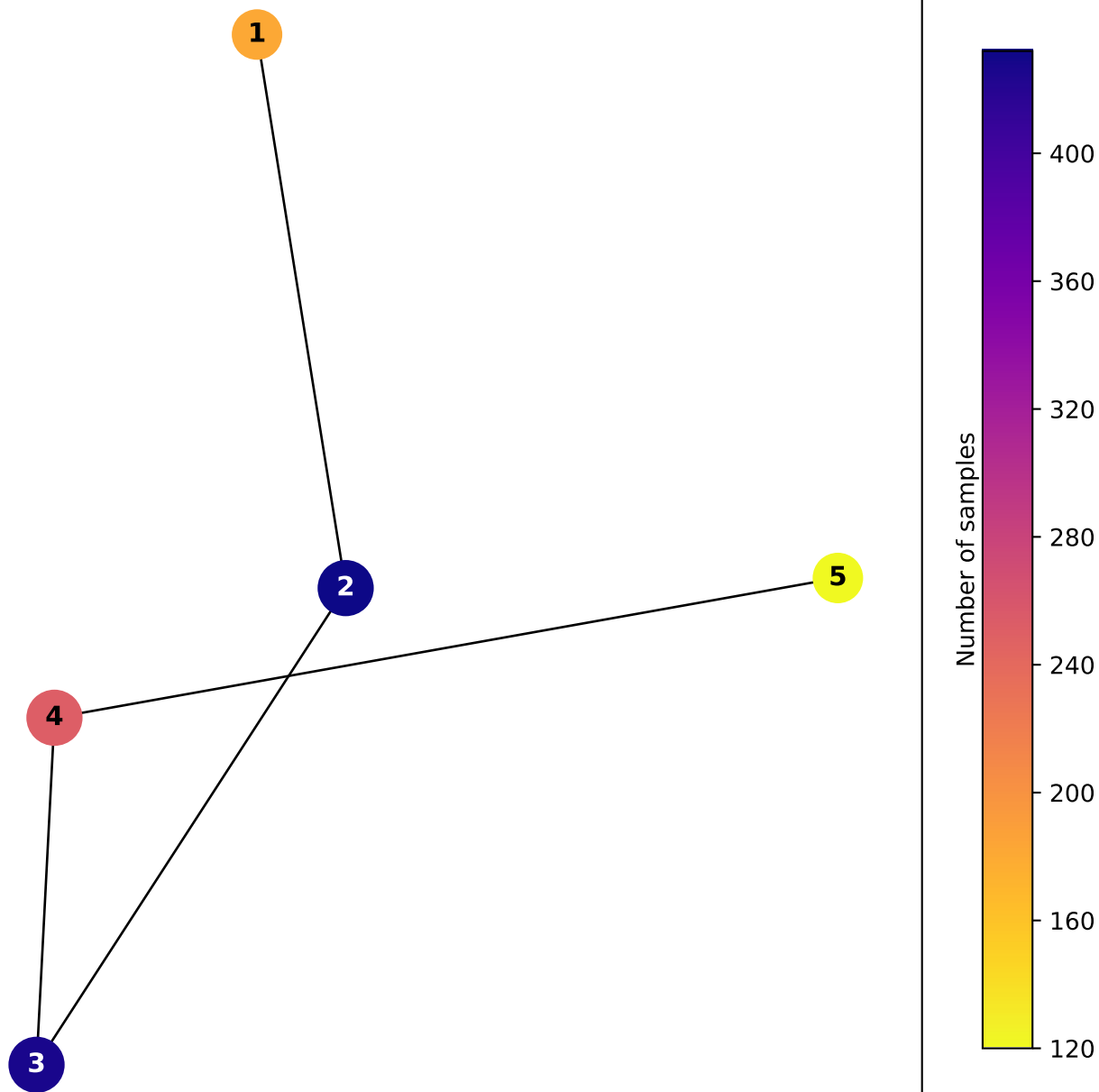
Intervals: 5 - Overlap: 0.25



Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

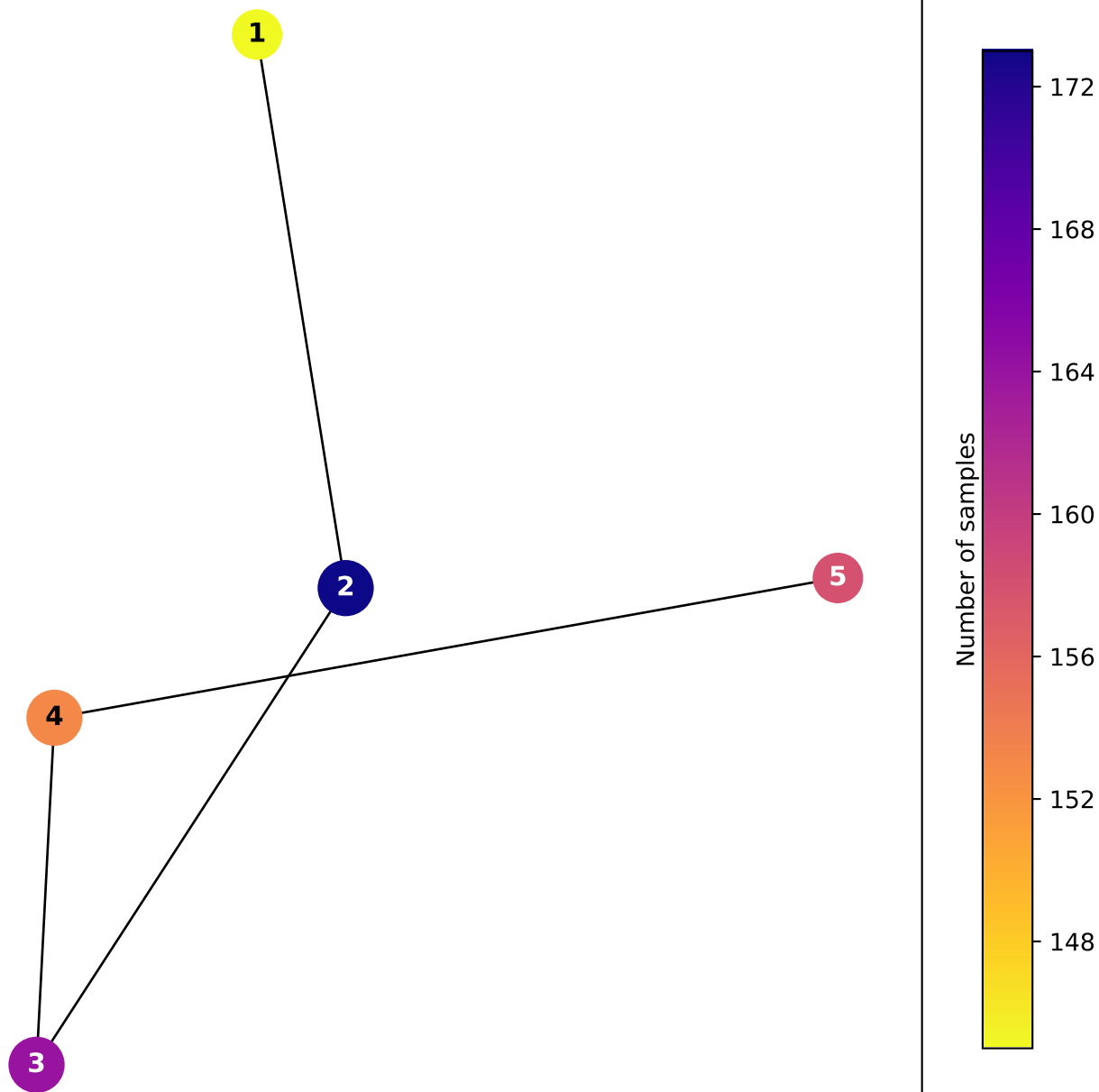
Intervals: 5 - Overlap: 0.5



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

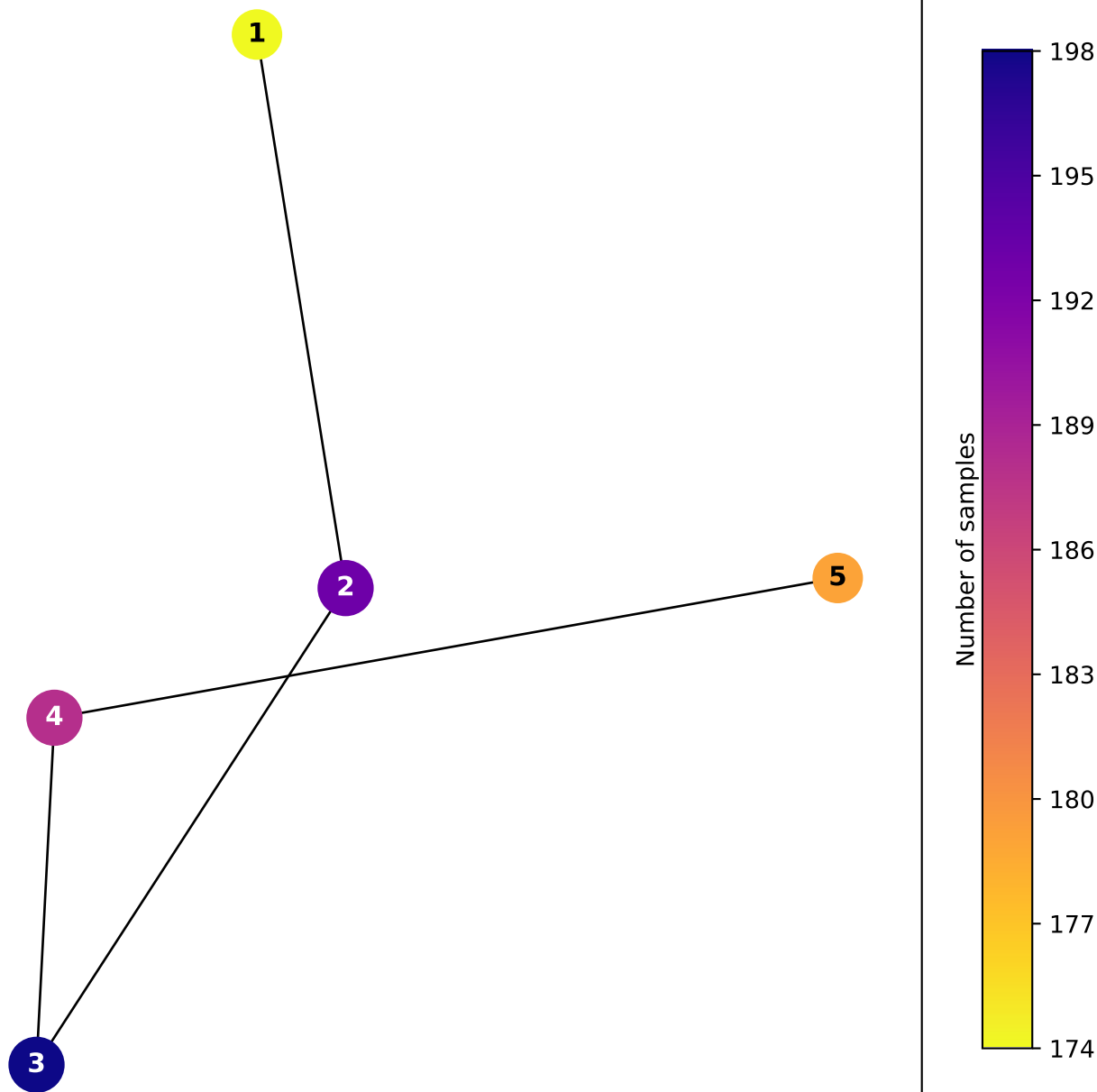
Intervals: 5 - Overlap: 0.1



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

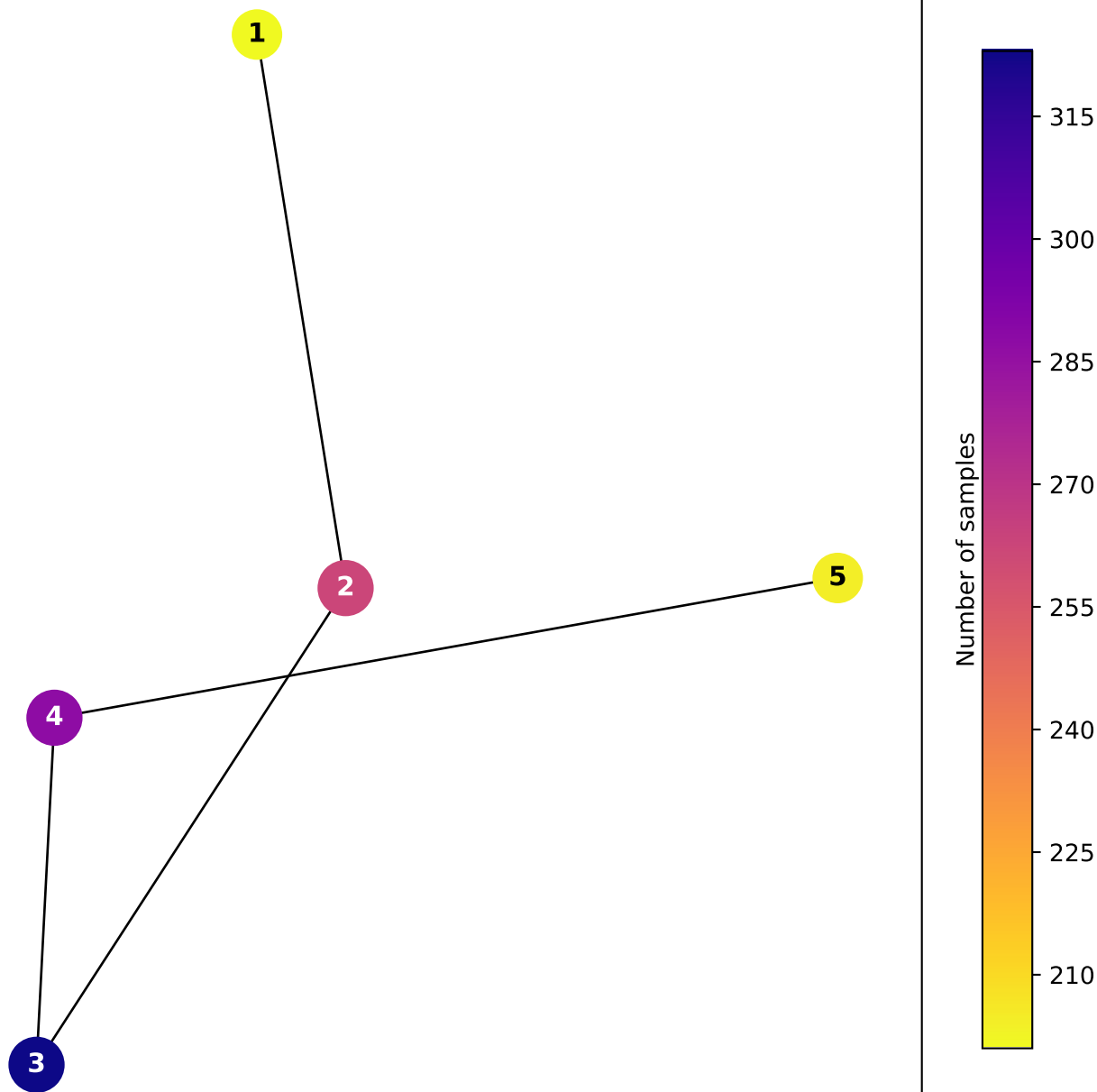
Intervals: 5 - Overlap: 0.25



Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)

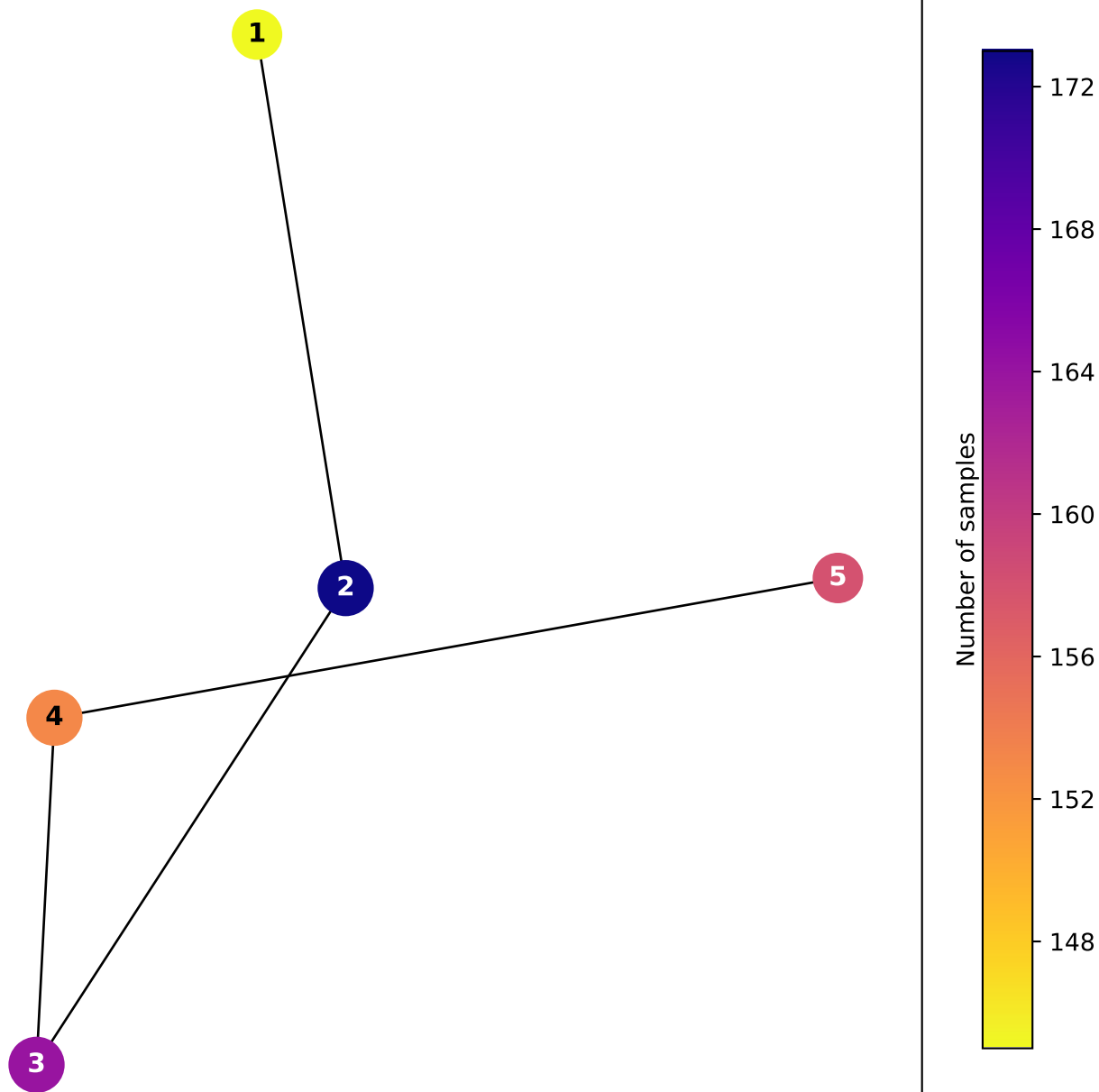
Intervals: 5 - Overlap: 0.5



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

Intervals: 5 - Overlap: 0.1



Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

Intervals: 5 - Overlap: 0.25

